

## SEQUENCE LISTING

SEQ ID NO: 1

Human unknown cDNA PHG-1 (GenBank # AL832747)

2110 bp

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1  gggatcgttc gatttaagcc atcatcagct taatttaagt tttagtttt tgctgaagga
61  ttatatgtat taatacttac ggttttaaat gtgttgcttt ggatacacac atagtttctt
121 ttttaataga atatactgtc ttgtctcact ttggactggg acagtggatg cccatctaaa
181 agttaagtg catttcttt agatgtttac cttcagccat agcttgattg ctcagagaaa
241 tatgcagaag gcaggatcaa agacacacag gagtcccttc ttttgaaatg ccacgtgcc
301 ttgtctttcc tcccttcttt gcttcttttt cttaccctct ctttcaattg cagatgccaa
361 aaaagatgcc aacagacact acattaccct aatggctgct acccagaacc ttttatagg
421 ttgtcttaa ttttttgtt gttgtgttc aagctttcc tttcttttt ttcttggtgt
481 ttgggccacg attttaaagt gacttttatt atgggtatgt gtgccaaag ctggctttt
541 gtcaaataaa atgaatacga acttaaaaaa taaaagctgg tatcttaaaa tgtaagagag
601 taagactgtg aagcctaaaa tgactggctg agaataaacc agaaatgcc tttgccaaac
661 agttgtaact agaaatttga ttctcaggtt ccattctttt ctttgcctt aagatgacat
721 ttgttagtgt cacgtcccat gttcagtgtc caaaccggca atgtaaaaag tatccttgt
781 ggtttaacag gaaatctgtt tatgtctctt tatttgaac cagttttact ctcagtgtt
841 ctttaagttc aatgaagtct gccaggaaca ttggttggtg gtattattcc gacaccttta
901 atttcaaaaa tctgaagttc ctgctagttt accaccttca tgatcttctt gaactggtaa
961 ctgattaggt tgaacttatg gaagatttgt ggacttaact caaaagtaac ctctcagtgt
1021 tctatagaac atgtatttgt gtaactgaac ctaccaggag aaatgtttgg aattctatat
1081 gtgcaatttt tcaacaaatg caaaaaaaat acagcacatg tattgacaag cttctgtcaa
1141 gcagcttgag ttgaaatttg atttaagaaa ataaatcatg attgttcaaa gctgctggga
1201 cgttagaatt aggccatgat actggctcct ttttaactac agtggtattt ggcactagt
1261 taaacttcca tataaatcac tcttttgtaa caacaaaggg ggaggaggaa aaatcacggc
1321 ctgttaaatg agtaccaaaag ccgccaaca gtaatgagat gttctcatcc ttgattctcc
1381 cagcctcaaa caacacagct tactttttt ttccctgtct cagaaagtac ctgtaattta
1441 acaaacagac tgctgttagg tatagtcaa ttacaaatgc tctaatcatt gtacatacat
1501 ctctcttgat attgcagcat ccatactggc ttgtaatca ttaattttt ggcagattga
1561 atgtgctgta ttgatatgta tctatgtaat tgtattgtat gtctatagct aattcacgtt
1621 ttgaataatg ttattttatt tactttttt agagaggaga atgtaaattt gtcagtttat
1681 ttctgactag ggatatttct ttccattta gaaaagaaga aaaaaaaaaa accttactgt
1741 catacagagc ggtactagcg tcgtgctgta taaaatcatt tgcacattcc tgagtagagg
1801 tatactgatt ataagacca aaggtaatth catagcaaaa tacataaaat cagtcggagc
1861 tttatacaa acatggaaac caactttgta gaacttttgc catttgatct aggattggaa
1921 tatgagcttt tatacaattc atattcttat ttggcaaatg cacagttag tattacctct
1981 ctgatggcct ttactagaaa ggcagtttta gaagctattg tgatccacta aggaaatgtt
2041 ttaacagcta gagaccactg ctgcctgaa agggcgttct taaatttggg gcagcaaaaa
2101 aaagaaaaaa

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SEQ ID NO: 2

Human prostaglandin D2 Synthase cDNA (GenBank# NM\_000954)

775 bp

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1 tgcaggagaa tggctactca tcacacgctg tggatgggac tggccctgct ggggggtgctg
61 ggcgacctgc aggcagcacc ggaggcccag gtctccgtgc agcccaactt ccagcaggac
121 aagttcctgg ggcgctggtt cagcgcgggc ctgcctcca actcgagctg gctccgggag
181 aagaaggcgg cgttgtccat gtgcaagtct gtggtggccc ctgccacgga tgggtggctc
241 aacctgacct ccaccttctt caggaaaaac cagtgtgaga cccgaaccat gctgctgcag
301 cccgcggggt cctcgggctc ctacagctac cggagtcccc actggggcag cacctactcc
361 gtgtcagtgg tggagaccga ctacgaccag tacgcgctgc tgtacagcca gggcagcaag
421 ggccctggcg aggacttccg catggccacc ctctacagcc gaaccagac cccagggct
481 gagttaaagg agaaattcac cgccttctgc aaggcccagg gcttcacaga ggataccatt
541 gtcttctgc cccaaaccga taagtgcatt acggaacaat aggactcccc agggctgaag
601 ctgggagccc ggccagccag gtgaccccca cgctctggat gtctctgctc tgttcttcc
661 ccgagcccct gcccgggctc cccgccaag caccctgcc cactcgggct tcactctgca
721 caataaactc cggaagcaag tcagttaaaa aaaaaaaaaa aaaaaaaaaa aaaaa

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SEQ ID NO: 3

Human myelin basic protein cDNA (GenBank# M13577)

2139 bp

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1 gaaaacagtg cagccacctc cgagagcctg gatgtgatgg cgtcacagaa gagaccctcc
61 cagaggcacg gatccaagta cctggccaca gcaagtacca tggaccatgc caggcatggc
121 ttctcccaa ggacagaga caggggcatc ctgactcca tcgggcgctt ctttggcggt
181 gacaggggtg cgccaaagcg gggctctggc aaggactcac accaccggc aagaactgt
241 cactatggct cctgcccc gaagtcacac ggccggaccc aagatgaaaa cccgtagtc
301 cacttcttca agaactgtg gacgcctcgc acaccacccc cgtcgcaggg aaaggggaga
361 ggactgtccc tgagcagatt tagctggggg gccgaaggcc agagaccagg atttggtac
421 ggaggcagag cgtccgacta taaatcggt cacaagggat tcaaggaggt cgatgccag
481 ggcacgcttt ccaaaatttt taagctggga ggaagagata gtcgctctgg atcacccatg
541 gctagacgct gaaaaccac ctggttccgg aatcctgtcc tcagcttctt aatataactg
601 ccttaaaact ttaatccac ttgccctgt tacctaatta gacagatga cccctccct
661 aatgcctgcg gagttgtgca cgtagtaggg tcaggccacg gcagcctacc ggcaatttcc
721 ggccaacagt taaatgagaa catgaaaaca gaaaacggtt aaaactgtcc ctttctgtg
781 gaagatcacg ttcttcccc cgcaatgtgc cccagacgc acgtgggtct tcaggggggc
841 aggtgcacag acgtccctcc acgttcaccc ctccaccctt ggactttctt ttcgccgtgg
901 ctgggcaccc ttgcgctttt gctggtcact gccatggagg cacacagctg cagagacaga
961 gaggacgtgg gcggcagaga ggactgttga catccaagct tcctttgttt ttttctctg
1021 tccttctctc acctcctaaa gtagacttca ttttcttaa caggattaga cagtcaagga
1081 gtggcttact acatgtggga gctttttgt atgtgacatg cgggctgggc agctgttaga
1141 gtccaacgtg gggcagcaca gagagggggc cacctcccca ggccgtggct gccacacac
1201 cccaattagc tgaattcgcg tgtggcagag ggaggaaaag gaggcaaacg tgggctgggc
1261 aatggcctca cataggaaac aggtcttcc tggagatttg gtgatggaga tgtcaagcag
1321 gtggcctctg gacgtcaccg ttgccctgca tgggtggccc agagcagcct ctatgaacaa
1381 cctcgtttcc aaaccacagc ccacagccgg agagtccagg aagacttgcg cactcagagc

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1441 agaagggtag gagtcctcta gacagcctcg cagccgcgcc agtcgccc at agacactggc  
 1501 tgtgaccggg cgtgctggca gggcagtg acagtggcca gactaacc tcctgagaa  
 1561 gataaccggc tcattcactt cctcccagaa gacgcgtggt agcgagtagg cacaggcgtg  
 1621 cacctgtcc cgaattact accgagacac acgggctgag cagacggccc ctgtgatgga  
 1681 gacaaagagc tcttctgacc atatccttct taacacccgc tggcatctcc ttctgcgcct  
 1741 ccctccctaa cctactgacc caccttttga ttttagcgca cctgtgattg ataggccttc  
 1801 caaagagtcc cagctggca tcacctccc cgaggacgga gatgaggagt agtcagcgtg  
 1861 atgcaaaac gcgtcttctt aatccaattc taattctgaa tgttctgtgt gggcttaata  
 1921 ccatgtctat taatatatag cctcgatgat gagagagtta caaagaacaa aactccagac  
 1981 acaaacctcc aaatttttca gcagaagcac tctgcgtcgc tgagctgagg tcggctctgc  
 2041 gatccatagc tggccgcacc cacacagcac gtgctgtgac gatggctgaa cggaaagtgt  
 2101 acactgttcc tgaatattga aataaaacaa taaactttt

SEQ ID NO: 4

Human unknown cDNA PHG-4 (GenBank# AP006241)

166 bp

1 ttcatataca aaaagataaa acttgaaata gttctagatt ttctctcta  
 51 ttgttgggt gtaactgctt cttcacacag ggggaaaaaa ctacattcac  
 101 atcggtttat ttgaggaccc agtcagagt tcaagcagca aaacccaac  
 151 ttagcagatc taattt

SEQ ID NO: 5

Human unknown cDNA PHG-5 (GenBank# BC011973)

1618 bp

1 ggcttggta ccgcattaag gcattccgc tctccgcgga actgctctgc cgtctcggcg  
 61 gtgaaagtgt gagagggtcc gtatgtgggt caacttgac tctctcgcc tgcccggatc  
 121 ctaagggcc tctcgtcct cccggtctcc ggtcgtgcc gggctctgac gccggtccgc  
 181 gcccggcctc gctctgcat gggcgctcc agtcctccg cgctggccc cctcggcctc  
 241 ccagccggc cctggcccag gtggctcggg gtcgccgcgc taggactggc cgcctggcc  
 301 ctggggactg tcgctggcg ccgcgcattg ccagggcggc gccggcggct gcagcaggtg  
 361 ggcaccgtgg cgaagctctg gatctaccg gtgaaatcct gcaaagggt gccggtgagc  
 421 gaggtgagt gcacggccat ggggtgcgc agcggcaacc tgcgggacag gtttggctg  
 481 gtgattaagg aagatggaca catggtcact gccgacagg agcctcgcct cgtgtctc  
 541 tccatcatt atgagaataa ctgcctgac ttacaggctc cagacatgga ccagctggtt  
 601 ttgctagca agcagccttc ctcaacaaa ctccacaact gcaggatatt tggccttgac  
 661 attaaaggca gagactgtgg caatgaggca gtaagtgtg taccactt ctgaaaact  
 721 gaagcgtata gattggttca attgagaca aacatgaagg gaagaacatc aagaaaact  
 781 ctcccactc ttgatcagaa ttccagggt gcctaccag actactgcc gctcctgac  
 841 atgacagatg cctccctggt agattgaat accaggatgg agaagaaaat gaaaatggag  
 901 aatttcaggc caaatattgt ggtgaccggc tgtgatgctt ttgaggagga tacctgggat  
 961 gaactcctaa ttgtagtgt agaagtgaag aaggtaatgg catgcccag gtgtattttg

1021 acaacggtgg acccagacac tggagtcata gacaggaaac agccactgga caccctgaag  
 1081 agctaccgcc tgtgtgatcc ttctgagagg gaattgtaca agttgtctcc actttttggg  
 1141 atctattatt cagtggaaaa aattggaagc ctgagagttg gtgaccctgt gtatcggatg  
 1201 gtgtagtat gatgatgga tccactaggg tgatatggct tcagcaacca ggagggattg  
 1261 actgagatct taacaacagc agcaacgata catcagcaaa tccttattat ccagccttca  
 1321 actatcttta ccctggaaaa caatctcgat tttgacttt tcaaagtgt gtatgtcca  
 1381 ggtaaatgca aggaaagtat tagagggggg aatatgaaag tatatatata aatttaggt  
 1441 actgaaggct ttaaaaataa ttaagatcat caaaaatgct atttgaatg ttatcatggc  
 1501 tattacactt ttacttctg actttaatat tgatgaataa agcaagtta atgaatcaac  
 1561 taaaagctg caaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaa

SEQ ID NO: 6

Human peanut-like 2/septin 4 cDNA (GenBank# NM\_080416)

1669 bp

1 cggcggtgct gcgaggtcgg cgcgcagctc cgccgcggtt cgctcgggcg ctgtccaggc  
 61 ggagccggcc ccgcccgggc tgcagccatg atcaagcgtt tcttgagga caccacggat  
 121 gatggagaac tgagcaagt cgtgaaggat ttctcaggaa atgcgagctg ccaccacca  
 181 gaggctaaga cctgggcac cagggcccaa gtcccggagc caaggcccca ggccccggac  
 241 ctctatgatg atgacctgga gttcagacc ccctcgcggc ccagtcctc tgacaaccag  
 301 cagtactct gtgcccagc ccctctcagc ccactgcca ggcccgcag cccatggggc  
 361 aagcttgatc cctatgattc ctctgaggat gacaaggagt atgtgggctt tgcaaccctc  
 421 cccaaccaag tccaccgaaa gtccgtgaag aaaggctttg actttaccct catggtggca  
 481 ggagagtctg gcctgggcaa atccacactt gtcaatagcc tcttctcac tgatctgtac  
 541 cgggaccgga aacttcttgg tgctgaagag aggatcatgc aaactgtgga gatcactaag  
 601 catgcagtgg acatagaaga gaagggtgtg aggctcggc tcaccattgt ggacacacca  
 661 ggttttggg atgcagtcaa caacacagag tgctggaagc ctgtggcaga atacattgat  
 721 cagcagttg agcagtattt ccgagacgag agtggcctga accgaaagaa catccaagac  
 781 aacagggtgc actgctgcct gtacttcac tcacccttcg gccatgggct ccggccattg  
 841 gatgttgaat tcatgaaggc cctgcatcag cgggtcaaca tcgtgcctat cctggctaag  
 901 gcagacacac tgacacctcc gaaagtggac cacaagaaac gaaaatccg ggaggagatt  
 961 gagcattttg gaatcaagat ctatcaatc ccagactgtg actctgatga ggatgaggac  
 1021 ttcaaatgac aggaccaagc cctaaaggaa agcatcccat ttgcagtaat tggcagcaac  
 1081 actgtagtag aggccagagg gcggcgagtt cggggtcgac tctaccctg gggcatcgtg  
 1141 gaagtggaaa acccagggca ctgcgacttt gtgaagctga ggacaatgct ggtacgtacc  
 1201 cacatgcagg acctgaagga tgtgacgagg gagacacatt atgagaacta ccgggcacag  
 1261 tgcattcaga gcatgaccg cctggtggtg aaggaaacgga atcgcaaaa actgactcgg  
 1321 gaaagtggta ccgacttccc catcctgct gtcccaccag ggacagatcc agaaactgag  
 1381 aagcttatcc gagagaaaga tgaggagctg cggcggtatg aggagatgct acacaaaata  
 1441 caaaaacaga tgaaggagaa ctattaactg gcttcagcc ctggatattt aaatctctc  
 1501 ctcttcttcc tgtccatgcc ggcccctccc agcaccagct ctgctcaggc cccttcagct  
 1561 actgccactt cgccttcat cctgtgtgac tgcccagaga ctgagaggaa ataaagtta  
 1621 ataatctgt aggtggctaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaa

SEQ ID NO: 7

Human coactosin-like 1 cDNA (GenBank# NM\_021149)

1850 bp

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1  cgcgctcgca gctcgcaggc gccgcgtagc cgtgccacc gccgccagcc cgtgcgcct
61  cggcgcgtac ccgcgcgct cccatccccg ccgccggcca ggggcgcgct cggccgccc
121 ggacagtgtc ccgctcggc tccgcggcga tggccaccaa gatcgacaaa gaggcttgcc
181 gggcggcgta caacctgtg cgcgacgacg gctcggcgt catctgggtg acttttaaat
241 atgacggctc caccatgct cccggcgagc agggagcggg gtaccagcac tcatccagc
301 agtgcacaga tgacgtcgg tgtttgcct tcgtgcgtt caccaccggg gatgccatga
361 gcaagaggtc caagtttgc ctcacacgt gcatcggta gaacgtcagc gggctgcagc
421 gcgcaaaaac cgggacggac aagaccctgg tgaaggaggt cgtacagaat ttcgtaagg
481 agtttgtgat cagtatcgg aaggagctgg aggaagattt catcaagagc gagctgaaga
541 aggcgggggg agccaattac gacgccaga cggagtaacc ccagccccg ccacaccacc
601 ccttgccaaa gtcatctgc tctccccg gggagaggac cgcggcctc agctactagc
661 ccaccagccc accagggaga aaagaagcca tgagaggcag cggccgccac cctgtgtcca
721 cagccccac ctcccgtt ccttagaac cctgccgtt cctatctcat gacgtcatg
781 gaacctctt cttgatctt cttttctt tctccccct tttttgtt taaagaaaag
841 tcattttgat gcaaggtcct gcctgccatc agatccgagg tgcctcctgc agtgaccctt
901 tttcttgga tttctctcc acgcgacgag gtctgcctag tgagatctgc atgacctac
961 gttgcttcc agagccccg cctatttgc catctcagtt ttctggacc ctgcttctg
1021 tgtaccactg aggggcagct gggccaggag ctgtgcccg tgcctgcagc cttcataagc
1081 acacacgtcc attccctact aaggccaga cctcctgga tctgcccg gctccctcat
1141 cccacctcca tccggagttg cctaagatgc atgtccagca taggcaggat tgctcggtgg
1201 tgagaaggtt aggtccggct cagactgaat aagaagagat aaaatttgc ttaaaactta
1261 cctggcagtg gctttgtgc acggtctgaa accacctgt cccacctct tgaccgaaat
1321 ttccttgta cacagagaag ggcaaagtc tgagccaga gttgacggag ggagtatttc
1381 agggttcact tcaggggctc ccaaagcgac aagatcgta gggagagagg cccagggtgg
1441 ggactgggaa ttaaggaga gctgggaacg gatcccttag gttcaggaag cttctgtgta
1501 agctgcgagg atggcttggg ccgaagggtt gctctgccg ccgcgctagc tgtgagctga
1561 gcaaagccct gggctcacag cccccaaaa gcctgtggct tcagtctgc gtctgcacca
1621 cacattcaaa aggatcgtt tgtttgtt ttaaagaaag gtgagattgg ctggttctt
1681 catgagcaca ttgatatag ctcttttct gttttcctt gtcatttcg tttggggaa
1741 gaaatctgta ctgtattggg attgtaaaga acatctctgc actcagacag ttacagaaa
1801 taaatgttt tttgtttt cagaaaaaaa aaaaaaaaaa aaaaaaaaaa

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SEQ ID NO: 8

Human clusterin mRNA (GenBank# BC019588)

1646 bp

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1  ctgaccgagg cgtgcaaaga ctccagaatt ggaggcatga tgaagactct gctgctgtt
61  gtggggctgc tgctgacctg ggagagtggg caggtcctgg gggaccagac ggtctcagac
121 aatgagctcc aggaaatgtc caatcaggga agtaagtacg tcaataagga aattcaaaat
181 gctgtcaacg ggggtgaaca gataaagact ctcatagaaa aaacaaacga agagcgcaag
241 aactgtctca gcaacctaga agaagccaag aagaagaaag aggatgccct aatgagacc
301 agggaatcag agacaaagct gaaggagctc ccaggagtgt gcaatgagac catgatggcc

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361 ctctgggaag agtgaagcc ctgcctgaaa cagacctgca tgaagttcta cgcacgcgtc  
421 tgcagaagtg gctcaggcct gggtggccgc cagcttgagg agttcctgaa ccagagctgc  
481 ccccttact tctggatgaa tggtagccgc atcgactccc tgctggagaa cgaccggcag  
541 cagacgcaca tgctggatgt catgcaggac cacttcagcc gcgcgtccag catcatagac  
601 gagctcttcc aggacaggtt ctaccccgagg gagccccagg atacctacca ctacctgccc  
661 ttcagcctgc cccaccggag gcctcacttc ttctttccca agtcccgcat cgtccgcagc  
721 ttgatgccct tctctccgta cgagcccctg aacttcacg ccatgttcca gcccttctt  
781 gagatgatac acgaggctca gcaggccatg gacatccact tccacagccc ggcttccag  
841 caccgccaa cagaattcat acgagaaggc gacgatgacc ggactgtgtg ccgggagatc  
901 cgccacaact ccacgggctg cctgcggatg aaggaccagt gtgacaagt ccgggagatc  
961 ttgtctgtg actgttccac caacaacccc tcccaggcta agctgcggcg ggagctcgac  
1021 gaatccctcc aggtcgctga gaggttgacc aggaaataca acgagctgct aaagtcctac  
1081 cagtgggaaga tgctcaacac ctctccttg ctggagcagc tgaacgagca gttaactgg  
1141 gtgtccggc tggcaaacct cagcaaggc gaagaccagt actatctgc ggtcaccacg  
1201 gtggcttccc acacttctga ctggacgtt ccttccgggtg tctactgaggt ggtcgtgaag  
1261 ctctttgact ctgatccat cactgtgacg gtccctgtag aagtctccag gaagaacct  
1321 aaatttatgg agaccgtggc ggagaaagcg ctgcaggaat accgcaaaa gcaccgggag  
1381 gactgagatg tggatgttc tttgcacct acgggggcat ctgagtccag ctcccccaa  
1441 gatgagctgc agccccccag agagagctct gcacgtcacc aagtaaccag gccccagcct  
1501 ccaggcccc aactccgcc agcctctccc cgctctggat cctgcactct aacactcgac  
1561 tctgtgctc atgggaagaa cagaattgct cctgcatgca actaattcaa taaaactgtc  
1621 ttgtgagctg aaaaaaaaaa aaaaaa

SEQ ID NO: 9

Human casein kinase 1, epsilon cDNA (GenBank# NM\_152221)

1559 bp

1 gggaggcggc ggcggcggcg gcggcggcgg cgagagccca gagccagagc ccggccgggg  
61 ccgagcggag cgcggcggcg gcggcggcgg cggcggctgg gccgggagag gctggcgcgc  
121 cgggcggctc cgcgaatct ccggcatccg ccccgcgagg ccgccccgc ccgcccagc  
181 ccccgagca gtggccggc atcggcgcct tcccgcgagg caagagttag ccatggagct  
241 acgtgtgggg aacaagtacc gcctgggacg gaagatcggg agcgggtcct tcggagatat  
301 ctacctgggt gccaatatcg cctctggtga ggaagtcgcc atcaagctgg agtgtgtgaa  
361 gacaaagcac cccagctgc acatcgagag caagttctac aagatgatgc aggggtggcgt  
421 ggggatcccc tccatcaagt ggtgcggagc tgagggcgac tacaactga tggatcatga  
481 gctgtgggg cctagcctcg aggacctgt caacttctgt tcccgcaaat tcagcctcaa  
541 gacggtgctg ctctggccg accagatgat cagccgcacg gattatatcc actccaagaa  
601 ctcatccac cgggacgtca agccgacaa ctctctcatg gggctgggga agaagggcaa  
661 cctgtctac atcatcgact tcggcctggc caagaagtac cgggacgccc gcaccacca  
721 gcacattccc taccgggaaa acaagaacct gaccggcacg gcccgtacg ctccatcaa  
781 cagcacctg ggcattgagc aaagccgtcg agatgacctg gagagcctgg gctacgtgct  
841 catgtacttc aacctgggt cctgccctg gcaggggctc aaagcagcca ccaagcgcca  
901 gaagtatgaa cggatcagcg agaagaagat gtcaacgcc atcgaggtcc tctgcaaagg  
961 ctatccctcc gaattctcaa catacctcaa ctctgccgc tcctgcgggt ttgacgaaa  
1021 gcccgactac tcttacctac gtcagctctt ccgcaacctc ttccaccggc agggcttctc  
1081 ctatgactac gctttgact ggaacatgct gaaattcggg gcagcccga atcccaggga

1141 tgtggaccgg gagcggcgag aacacgaacg cgaggagagg atggggcagc tacgggggtc  
 1201 cgcgaccga gccctgcccc ctggcccacc cacggggggcc actgccaacc ggctccgcag  
 1261 tgccgccgag cccgtggctt ccacgccage ctcccgcatc cagccggctg gcaatacttc  
 1321 tcccagagcg atctcgcggg tcgaccggga gaggaaggtg agtatgaggc tgcacagggg  
 1381 tgcgcccgc aacgtctct cctcagacct cactgggcgg caagaggctt cccggatccc  
 1441 agcctcacag acaagtgtgc catttgacca tctcgggaag tgaggagagc cccattgga  
 1501 ccagtgtttg cttagtgtct tctgttatt ttcttataaa aaaaaaaaaa aaaaaaaaaa

SEQ ID NO: 10

Human ferritin, heavy polypeptide 1 cDNA (GenBank# BC015946)

910 bp

1 cctgcttcaa cagtgttgg acggaacccg gcgtcgttc cccaccccgg ccggccgccc  
 61 atagccagcc ctccgtacc tcttcaccg accctcggac tgcccaagg ccccgccgc  
 121 cgctccagcg ccgcgcagcc accgcccgcg ccgcccctc tcttagtcg ccgccatgac  
 181 gaccgcgtcc acctgcagg tgcgccagaa ctaccaccag gactcagagg ccgccatcaa  
 241 ccgcccagat aacctggagc tctacgcctc ctacgtttac ctgtccatgt ctactactt  
 301 tgaccgcgat gatgtggtt tgaagaactt tgccaaatac ttcttcacc aatctcatga  
 361 ggagagggaa catgttgaga aactgatgaa gctgcagaac caacgaggtg gccgaatctt  
 421 ccttcaggat atcaagaaac cagactgtga tgactgggag agcgggctga atgcaatgga  
 481 gtgtgcatta catttgaaa aaatgtgaat cagtcactac tggaactgca caaactggcc  
 541 actgacaaaa atgaccccca ttgtgtgac ttcatlgaga cacattacct gaatgagcag  
 601 gtgaaagcca tcaaagaatt gggtgaccac gtgaccaact tgcgcaagat gggagcggcc  
 661 gaatctggct tggcgggaata tctctttgac aagcacaccc tgggagacag tgataatgaa  
 721 agctaagcct cgggctaatt tccccatagc cgtgggggtga ctccctggt caccaaggca  
 781 gtgcatgcat gttgggggtt cctttacctt ttctataagt tgtacaaaa catccactta  
 841 agttcttga ttgtaccat tcttcaaat aaagaaattt ggtacccaaa aaaaaaaaaa  
 901 aaaaaaaaaa

SEQ ID NO: 11

Human metargidin cDNA (GenBank# NM\_003815)

2740 bp

1 cgctgccatg cggctggcgc tgctctgggc cctggggctc ctgggcgcgg gcagccctct  
 61 gccttcttgg ccgctcccaa atataggtgg cactgaggag cagcaggcag agtcagagaa  
 121 ggccccgagg gagcccttgg agccccaggt ccttcaggac gatctcccaa ttgcctcaa  
 181 aaaggtgctt cagaccagtc tgcctgagcc cctgaggatc aagtggagc tggacggtga  
 241 cagtcatact ctggagctgc tacagaatag ggagttggtc ccaggccgcc caacctggt  
 301 gtggtaccag cccgatggca ctccgggtgt cagtgaggga cacactttgg agaactgctg  
 361 ctaccaggga agagtgcggg gatatgcagg ctccctgggtg tccatctgca cctgctctgg  
 421 gctcagaggc ttgtgtgtcc tgacccaga gagaagctat acctggagc aggggcctgg  
 481 ggaccttcag ggtctccca ttatttcgag aatccaagat ctccacctgc caggccacac  
 541 ctgtgccctg agctggcggg aatctgtaca cactcagacg ccaccagagc acccctggg  
 601 acagcgccac attcgccgga ggcgggatgt ggtaacagag accaagactg tggagtgtg  
 661 gattgtggt gatcactcg aggccagaa ataccgggac ttccagcacc tgctaaaccg  
 721 cacactggaa gtggccctct tgctggacac attctccgg cccctgaatg tacgagtggc

781 actagtgggc ctggaggcct ggacccagcg tgacctggtg gagatcagcc caaacccagc  
 841 tgcaccctc gaaaacttc tccactggcg cagggcacat ttgtgcctc gattgcccc  
 901 tgacagtgcc cagctggtga ctggtacttc attctctggg cctacggtgg gcatggccat  
 961 tcagaactcc atctgttctc ctgacttctc aggaggtgtg aacatggacc actccaccag  
 1021 catcctggga gtgcctcct ccatagccca tgagttgggc cacagcctgg gcctggacca  
 1081 tgatttgctt gggaatagct gccctgtcc aggtccagcc ccagccaaga cctgcatcat  
 1141 ggaggcctcc acagacttcc taccaggcct gaacttcagc aactgcagcc gacgggacct  
 1201 ggagaaagcc ctctggatg gaatgggcag ctgcctcttc gaacggctgc ctacgtacc  
 1261 cccatggtc gctttctcg gaaatatgt ttggagccg ggcgagcagt gtgactgtg  
 1321 ctctctggat gactgcgtcg atccctgtg tgattcttg acctgccagc tgaggccagg  
 1381 tgcacagtgt gcatctgacg gacctgttg tcaaaattgc cagctgcgc cgtctggctg  
 1441 gcagtgtcgt cctaccagag gggattgtga ctgcctgaa ttctgccag gagacagctc  
 1501 ccagtgtccc cctgatgtca gcctagggga tggcgagccc tgcgtggcg ggcaagctgt  
 1561 gtgcatgcac gggcgttggt cctctatgc ccagcagtgc cagtcacttt ggggacctgg  
 1621 agcccagccc gctgcgccac ttgcctcca gacagctaat actcggggaa atgcttttgg  
 1681 gagctgtggg cgcaacccca gtggcagtta tgtgtcctgc acccctagag atgccattg  
 1741 tgggcagctc cagtgccaga caggtaggac ccagcctctg ctgggctcca tccgggatct  
 1801 actctgggag acaatagatg tgaatgggac tgagctgaac tgcagctggg tgcacctgga  
 1861 cctgggcagt gatgtggccc agcccctct gactctgcct ggcacagcct gtggccctgg  
 1921 cctggtgtgt atagaccatc gatgccagcg tgtggatctc ctgggggcac aggaatgtcg  
 1981 aagcaaatgc catggacatg gggctctgta cagcaacagg cactgctact gtgaggaggg  
 2041 ctgggcaccc cctgactgca ccactcagct caaagcaacc agtccctga ccacagggtc  
 2101 gctcctcagc ctctggtct tattggtcct ggtgatgctt ggtgccggct actggtaccg  
 2161 tgcccgcctg caccagcgac tctgccagct caagggaccc acctgccagt acagggcagc  
 2221 ccaatctggt cctctgaac ggccaggacc tccgcagagg gccctgtctg cacgaggcac  
 2281 taagtctcag gggccagcca agccccacc cccaaggaag ccaactgcctg ccgaccccca  
 2341 gggccgggtg ccatcgggtg acctgcccgg ccagggggt ggaatccgc ccctagtgtg  
 2401 accctccaga ccagcgccac cgctccgac agtgcctcgt ctctacctt gacctctcg  
 2461 gaggttccgc tgcctccaag ccggacttag ggcttcaaga ggcgggcgtg cctctggag  
 2521 tccctacca tgactgaagg cgccagagac tggcggtgtc ttaagactcc gggcaccgcc  
 2581 acgcgtgtc aagcaacact ctgcggacct gccggcgtag ttgcagcggg ggcttgggga  
 2641 ggggttgggg gttggacggg attgaggaag gtccgcacag cctgtctctg ctgattgca  
 2701 ataacgtga catctggga gcgttaaaaa aaaaaaaaaa

SEQ ID NO: 12

Human unknown cDNA PHG-13 (GenBank# AK026351)

1476 bp

1 gtttaatagc ttgaggaagg gagactttaa aaggacgtgt gtgagtgaag taggatatag  
 61 ccattaccac ggtgccagga cctgacagcg ttccaattct tttgcagca tggggaatca  
 121 aaggtggcat gccaagtca actcagggtc gaggtatcca cattgtccac atcaggcaag  
 181 cctgcactg acggttgagc ctcatggaga ggagcatgtg ttggaagag atccctttgt  
 241 taactgtttt gtggtgttct cttaaatgaa ttagagctca tgcccccttt ctggctttgc  
 301 tgttgatttt gtaggttaga gaatattcct gagagccttc ctttggccc ccagcttatg  
 361 ccaccactc tcttctctg gttgaattct ctgaaggaaa ggttcattgt ctattgtcct  
 421 gttagtcaat agtcttcata tataattgtg ttacatatat tgctgtagac tctcagaaat

481 cagggtagag ctttccctt gagcagtta atgagtgaat tcagcagcaa agtcgcaaga  
 541 aatggttctc cagccaggag aggttatgtt taccctctga ttgcccggtt tctctgcaca  
 601 cagtgatata gtattcagt agaggtgctg ttggcaccca gcagcacctt gggcacacag  
 661 catttcattg catgtcacag tgtacaagct accctctaata tcagaaagaa gagcattttg  
 721 cacagagaaa aataaaaaga tccatgaatg tcatcttita tcttttattt tcagttggct  
 781 gatgttgga ttttgttct tgcataaac ttgtaaacca atcttgccaa gatacaagtt  
 841 gttttggtt ttactacaa tgacctctg ttctctctgt ctgactgct gacgttctc  
 901 aatgattcta ttgtctattt tatgggaagc agccttccca taggtttcct ttacacact  
 961 gcagggttat ctttatactt taataaaaaa aaaaaaaa aaaaggacaa gaactgtcac  
 1021 taacctcatg gaggggttg cgtaaaacca tttagccac cttagacaa gggtagattc  
 1081 cgtgtgttt tttaagctc actgtaataa aatagatcta attcagcatt attgtgctac  
 1141 ctcaaaggta aaaaatgtt taaggcttc ttttggtcct gatttctata tacagtgtt  
 1201 gaaatgtctt tcatttgga ttattttta aattcttga gtgaattta tttaattc  
 1261 tttaattctt gtattttaa ctcagaagaa taagtattg aaactgac aattcttgc  
 1321 ctgtgtgtt aaacataaa tgaacagta ttaagaatta agtactgtt tgccataaac  
 1381 aaggttgatg ttcttttgt tgtgttaag gaaaccctag ggctcggctt tactctgat  
 1441 taataaaggc tgacaaatca aaaaaaaa aaaaaa

SEQ ID NO: 13

Human retinaldehyde binding protein 1 cDNA (GenBank# NM\_000326)

1679 bp

1 ggcacgaggt agagctccag gacattcagg taccaggtag cccaaggag gagctgccga  
 61 cctggcaggg aacaaccaag actggggta aatctcacag cctgcaagt gaagagaaga  
 121 actgaaccc aggtccaact ttgcccac agcaggctgc ctcttggtc tgacaggaag  
 181 tcacaactg gcctgactt cctatcctag ggaaggggccc ggctggagag gccaggacag  
 241 agaaagcaga tccctcttt ttccaaggac tctgtgtct ccataggcaa catgcagaa  
 301 ggggtgggca cgttcgcat ggtacctgaa gaggaacagg agtccgtgc ccaactggag  
 361 cagtcacaa ccaaggacca tggacctgc ttggcccg gcagccagct gccccgccac  
 421 acctgcaga agccaagga tgagctgaac gagagagagg agaccggga ggaggcagt  
 481 cgagagctgc aggagatggt gcaggcgag gcggcctcg gggaggagct ggcggtgcc  
 541 gtggcgga ggggtcaaga gaaggacagc ggcttctcc tgcgttcat ccgcgcacgg  
 601 aagttaacg tgggccgtgc ctatgagctg ctacagaggt atgtgaatt ccggctgcag  
 661 taccctgagc tcttgacag cctgtccca gaggtgtcc gctgcacat tgaagtggc  
 721 taccctgtg tctctctag tgggacaag tatggccgag tggctatgct ctcaacatt  
 781 gagaactggc aaagtcaaga aatcacctt gatgagatc tgcaggcata ttgcttcat  
 841 ctggagaagc tctggagaa tgaggaaact caaatcaat gcttctgcat cattgagaac  
 901 ttcaagggt ttacatgca gcaggctgt agtctccga ctacagatc caggaagatg  
 961 gtggacatgc tccaggattc ctccagcc cggttcaag ccatcactt catccaccag  
 1021 ccatggtact tcaccacgac ctacaatgtg gtcaagccct tctgaagag caagctgctt  
 1081 gagagggtct ttgtccagg gtagacatt tctgtttct accaggagat ctagagaaac  
 1141 atctgccct ctgactcgg gggcacgtg ccaagtatg atggcaaggc cgttgctgag  
 1201 cagctcttg gccccaggc ccaagctgag aacacagcct tctgaaaaca tctctgcca  
 1261 gctgaactgt agttagaat tctgggctc tctcaactg tctggaccc aaggctagga  
 1321 aagggtgct tgagatgact gtgtccccc cttagactcc ctaagcccga gtgagctcag  
 1381 gtgtaccct gttctcaat tgggggatg ggaataaagg agggggaatt ccttgaaca

1441 agaagaactg gggatagtta tattccacc tgccttgaa gctttaagac agtgatttt  
 1501 gtgtaaggtt gtattcaaa gactcgaatt cattttctca atcatttctt ttgtaacaga  
 1561 gttttacgac ttagagtctg tgaaaacagg caaggagccc gggttaaaat atccccctat  
 1621 tcgcccccaa aatgcaataa aagaagataa aagagagagg aaaaaaaaaa aaaaaaaaaa

SEQ ID NO: 14

Human actin, gamma 1 cDNA (GenBank# BC009848)

1962 bp

1 agctctcgca ctctgttctt ccgccgctcc gccgtcgcgt ttctctgccg gtcgcaatgg  
 61 aagaagagat cgccgcgctg gtcattgaca atggctccgg catgtgcaa gctggtttg  
 121 ctggggacga cgctccccga gccgtgttcc ctccatcgt cgggcgcccc agacaccagg  
 181 gcgtcatggt gggcatgggc cagaaggact cctacgtggg cgacgaggcc cagagcaagc  
 241 gtggcatcct gacctgaag taccctattg agcatggcat cgtcaccaac tgggacgaca  
 301 tggagaagat ctggcaccac accttctaca acgagctgcg cgtggccccg gaggagcacc  
 361 cagtgtgct gaccgaggcc cccctgaacc ccaaggccaa cagagagaag atgactcaga  
 421 ttatgttga gacctcaac accccggcca tgtacgtggc catccaggcc gtgctgtccc  
 481 tctacgcctc tgggcgcacc actggcattg tcatggactc tggagacggg gtcaccacaca  
 541 cgggtgccc atacgagggc tacgccctcc ccacgccat cctgcgtctg gacctggctg  
 601 gccgggacct gaccgactac ctcatgaaga tctcactga gcgaggctac agcttcacca  
 661 ccacggccga gcgggaaatc gtgcgcgaca tcaaggagaa gctgtgctac gtcgccctgg  
 721 acttcgagca ggagatggcc accgccgat cctcctcttc tctggagaag agctacgagc  
 781 tgcccgatgg ccaggtcatc accattggca atgagcgggt ccggtgtccg gaggcgctgt  
 841 tccagccttc ctctctgggt atggaatctt gcggcatcca cgagaccacc ttcaactcca  
 901 tcatgaagtg tgacgtggac atccgcaaag acctgtacgc caacacggtg ctgtcgggcg  
 961 gcaccacat gtaccgggc attgccgaca ggatgcagaa ggagatcacc gccttgccgc  
 1021 ccagcaccat gaagatcaag atcatgcac cccagagcg caagtactcg gtgtggatcg  
 1081 gtggctccat cctggcctca ctgtccacct tccagcagat gtggattagc aagcaggagt  
 1141 acgacgagtc gggccccctc atcgtccacc gcaaatgctt ctaaaccggac tcagcagatg  
 1201 cgtagcattt gctgcatggg ttaattgaga atagaaattt gccctggca aatgcacaca  
 1261 cctcatgcta gcctcacgaa actggaataa gccttcgaaa agaaattgtc ctgaaagctt  
 1321 gtatctgata tcagcactgg attgtagaac ttgtgtctga tttgacctt gtattgaagt  
 1381 taactgttcc ctttggattt tgtttaatac cctgtacata tctttgagt caaccttag  
 1441 tacgtgtggc ttgtcactt cgtggctaag gtaagaacgt gcttgtggaa gacaagtctg  
 1501 tggcttggtg agtctgtgtg gccagcagcc tctgatctgt gcagggtatt aacgtgtcag  
 1561 ggctgagtgt tctgggattt ctctagaggc tggcaagaac cagttgtttt gtcttgccgg  
 1621 tctgtcaggg ttggaagtc caagccgtag gaccagttt ctttcttag ctgatgtctt  
 1681 tggccagaac accgtgggct gttacttgc ttgagttgga agcggtttgc atttacgcct  
 1741 gtaaatgtat tcattcttaa ttatgtaag gtttttttg tacgcaattc tcgattcttt  
 1801 gaagagatga caacaaattt tggttttcta ctgttatgtg agaacattag gccccagcaa  
 1861 cacgtcattg tgtaaggaaa aataaaagt ctgccgtaac caaaaaaaaa aaaaaaaaaa  
 1921 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aa

SEQ ID NO: 15

Human matrix metalloproteinase, membrane associated, cDNA (GenBank# X83535)  
2365 bp

```

1  gaattcaagt tcagtccta ccgaagacaa aggcgccccg agggagtggc ggtgcgaccc
61  cagggcgtgg gcccggccgc ggagcccaca ctgcccggct gacccggtgg tctcggacca
121  tgtctccgc cccaagaccc tcccgtgtc tctgtctcc cctgtcacg ctccgcaccg
181  cgctcgctc cctcggtcg gcccaaagca gcagcttcag ccccgaaacc tggctacagc
241  aatatggcta cctgcctccc ggggacctac gtaccacac acagcgctca cccagtcac
301  tctcagcggc catcgtgcc atgcagaagt ttacggctt gcaagtaaca ggcaaagctg
361  atgcagacac catgaaggcc atgaggcgcc ccgatgtgg tgtccagac aagtttgggg
421  ctgagatcaa ggccaatgtt cgaaggaagc gctacgcat ccagggtctc aaatggcaac
481  ataatgaaat cactttctgc atccagaatt acaccccaa ggtgggcgag tatgccacat
541  acgaggccat tcgaaggcg ttccgcgtgt gggagagtgc cacaccactg cgttccgcg
601  aggtgcccta tgcctacatc cgtgagggcc atgagaagca ggccgacatc atgatctct
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721  atgcctactt cccaggcccc aacattggag gagacacca cttgactct gccgagcctt
781  ggactgtcag gaatgaggt atgaatgaa atgacatct cctgggtggt gtgcacgagc
841  tgggcatgc cctggggctc gagcattcca gtgaccctc ggccatcatg gcacccttt
901  accagtggat ggacacggag aattttgtgc tgccgatga tgaccgccgg ggcattccagc
961  aactttatgg ggtgagtc ggttcccca ccaagatgc cctcaacc aggactacct
1021  cccggccttc tgttctgat aaacccaaaa accccaccta tgggccaac atctgtgacg
1081  ggaacttga caccgtggc atgctccgag gggagatgtt tgtctcaag gagcgctggt
1141  tctggcgggt gaggaataac caagtgatgg atggataccc aatgccatt ggccagtct
1201  ggcggggcct gcctgcgtcc atcaacactg cctacgagag gaaggatggc aaattcgtct
1261  tctcaaaag agacaagcat tgggtgttg atgaggcgtc cctggaacct ggctacccca
1321  agcacattaa ggagctgggc cgagggtgc ctaccgaca gattgatgct gctctctct
1381  ggatgcccaa tggaaagacc tacttctcc gtggaacaa gtactaccgt ttcaacgaag
1441  agtcagggc agtggatagc gagtaccaca agaacaacaa agtctgggaa gggatccctg
1501  agtctcccag aggtcattc atgggcagcg atgaagtct cacttactc tacaagggga
1561  acaatactg gaaattcaac aaccagaagc tgaaggtaga accgggctac cccaagtcag
1621  ccttgaggga ctggatgggc tgccatcgg gagggcgcc cgatgagggg actgaggagg
1681  agacggaggt gatcatcatt gaggtggacg aggagggcg cgggcggtg agcgctgctg
1741  ccgtggtgct gccgtgctg ctgctgtcc tgggtgtggc ggtgggacta gcagtctct
1801  tctcagacg ccatgggacc cccaggcgac tctctactg ccagcgttcc ctgctggaca
1861  aggtctgacg cccaccgccc gcccggccac tctaccaca aggactttgc ctctgaagac
1921  cagtgtcagc aagtggtgg tgggtgggct gctccatcc gtccggagcc cctccccgc
1981  agcctcctg cttctcag tcccgtggt ggctcctc accctaccg cctgtagctt
2041  gtgtctgtcc agccccatct gaatgtgtg ggggtctgc acttgaaggc aggaccctca
2101  gacctcgctg gtaaaggta aatgggtgta tctgtcctt ttccatcccc tgacatact
2161  taacctctga actctgacct caggaggctc tgggcactcc agcctgaaa gcccgaagt
2221  taccagttg gcagcctccc gtcactctga ctaaaagaa tctcagagt gcatatttg
2281  aggtggaaag attgttcagt taccctaaag acttgaaag aaagaaagaa agaaagaaa
2341  aaaaaaaaaa aaaaaaaaaa aaaaa

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SEQ ID NO: 16

Human SWI/SNF related/OSA-1 nuclear cDNA. (GenBank# NM\_006015)

8595 bp

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1 aaagcggaga gtcacagcgg gccagggccc tggggagcgg agcctccacc gccccctca
61 ttccaggca agggcttggg gggaatgagc cgggagagcc gggtcccgag cctacagagc
121 cgggagcagc tgagccgccc gcgcctcggc cggcggccc gcctctctct cctccgccgc
181 cggcagcccc gagcctgagc cggcggggcg ggggggagag gagcagcgc agcgcagcag
241 cggagccccg cgaggcccg ccggcggggt ggggaggcca gcccggggga ctgggccccg
301 gggcggggtg ggaggggggg agaagacgaa gacagggccg ggtctctccg cggacgagac
361 agcggggatc atggccgcgc aggtcggccc cggcggccc agcagcctgg gcaaccgcc
421 gccggcccg ccctcggagc tgaagaaagc cgagcagcag cagcgggagg aggcgggggg
481 cgaggcggcg gcggcggcag cggccgagcg cggggaaatg aaggcagccg ccgggcagga
541 aagcaggggc cccgccgtgg ggccgccga gccgctggga aaggagctgc aggcgggggc
601 cgagagcaat ggggggtggc gcggcggcgg agccggcagc ggcgggggc ccggcgcgga
661 gccggacctg aagaactga acgggaacgc gggccctagg cccgccctga acaataacct
721 cacggagccg cccggcgcg gcggtggcgg cagcagcgat ggggtggggg cgcctctca
781 ctacgccg gcgccttg cggcccccag ctacggctt cggcaacct acggccggag
841 cccgtctgcc gtcgcccg ccgcgccgc cgtctccac caacaacatg gcggacaaca
901 aagccctggc ctggcagcgc tgcagagcgg cggcgggcgg gccctggagc cctacgcggg
961 gcccagcag aactctcag accacggct cccaaccac cagtacaact cctactacc
1021 caaccgcagc gcctacccc cggcgccc ggctacgcg ctgagctccc cgagaggtgg
1081 cactccgggc tccggcgcg gcggcgctgc cggctccaag ccgcctccct cctccagcgc
1141 ctccgctcc tctgtctt cgtcttcgc tcagcagcgc ttcggggcca tggggggagg
1201 cgccccctcc gcggccggcg ggggaactcc ccagcccacc gccaccccc cctcaacca
1261 actgctcag tcgcccag ctgcccgggg ctaccagggc taccggggg gcgactacag
1321 tggcgggccc caggacgggg gcgcggcga gggccggcg gacatggcct cgagtggtg
1381 gggggctgcg gcggcggcag ctgcggcggc gccgcctcg ggagggggcc acaaaggag
1441 ccaccagcgc ccatgagcc cgggagcag cggcgcggg gggcagccgc tcgcccggac
1501 cctcagcca tccagtcaa tggatcagat gggcaagatg agacctcagc catatggcgg
1561 gactaacca tactgcagc aacagggacc tccgtcagga ccgcagcaag gacatgggta
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1681 ggcgcagagt gccatggcg gcctcttta tacacagcag attctctt atggacaaca
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1801 cctcagcag cagcagccac cctactcca gcaaccaccg tccagaccc ctcatgccc
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2341 gagtctgga gtgagcacat cagggatct cagcagccaa ggagagcaga gtaatccagc
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2521 tgcagtgcc a ggcaaccaga tggcacctcg gccacccagt ggccagtcgg acagcatcat  
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SEQ ID NO: 17

Human unknown cDNA AMDP-3 (GenBank# AK024103)

3488 bp

1 taaaaagcat taggcatata aatgtataaa tatatttat catgtacagt aaaaaatgg  
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 121 agttcttgc gatgcttgca cctgacagt gggaccaac acagacgtgc cacccaaccc  
 181 cctgcacaca ccaccggcca ccagggggccc ccttgctgcg cttggcttta taactcctc  
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 3361 gaaatagaga tgcagtactt aactttcctt ggtgtttgta gatattgcct tgtgtattcc  
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SEQ ID NO: 18

Human MT1-MMP exon 1s

20 bp

5'GCCTACCGAAGACAAAGGCG3'

SEQ ID NO:19

Human MT1-MMP exon 1a

20 bp

5'TAGAGGCTGTCCCCTAGGAG3'

SEQ ID NO:20

Human MT1-MMP exon 2s

20 bp

5'AGAGGCACCCTATGGGCCAG3'

SEQ ID NO:21

Human MT1-MMP exon 2a

20bp

5'CATCTCTGGCGCTGGCATTG3'

SEQ ID NO:22

Human MT1-MMP exon 3s

20 bp

5'GCACTGATCCCAATCCTCGC3'

SEQ ID NO:23

Human MT1-MMP exon 3a

20 bp

5'CCCTGCATAAGCACAATGGG3'

SEQ ID NO:24

Human MT1-MMP exon 4s

20 bp

5'GGGAAGGAGAATGTTGCCCC3'

SEQ ID NO:25

Human MT1-MMP exon 4a

20 bp

5'GAGGAGGGAACCACCCCTAC3'

SEQ ID NO:26

Human MT1-MMP exon 5s

20 bp

5'GGGAGGCTGAGGGAAGGGAC3'

SEQ ID NO:27

Human MT1-MMP exon 5a

20 bp

5'GGGGAAATGCGTAGACCAGG3'

SEQ ID NO:28

Human MT1-MMP exon 6s

20 bp

5'CCCGCCTCCTCCTAAGTCTG3'

SEQ ID NO:29

Human MT1-MMP exon 6a

20 bp

5'CAGCATGAGCCACCATGCCC3'

SEQ ID NO:30

Human MT1-MMP exon 7s

20 bp

5'GAACCAGAGACCTAGGCCGC3'

SEQ ID NO:31

Human MT1-MMP exon 7a

20 bp

5'CAGCTCCTCTAGGGAGACCC3'

SEQ ID NO:32

Human MT1-MMP exon 8s

20 bp

5'CTAGAGCCTAAGTTGAACCC3'

SEQ ID NO:33

Human MT1-MMP exon 8a

20 bp

5'GTGGTGGTGGTTTATGAGGG3'

SEQ ID NO:34

Human MT1-MMP exon 9s

20 bp

5'TAGGACATGCCCATGTCCGC3'

SEQ ID NO:35

Human MT1-MMP exon 9a

20 bp

5'TCCGCTCTTCCTCAACTCCC3'

SEQ ID NO:36

Human MT1-MMP exon 10s

20 bp

5'CTCTTTGGGTCTTCCTTCC3'

SEQ ID NO:37

Human MT1-MMP exon 10a

20 bp

5'CTTCAGAGGCAAAGTCCTTG3'

SEQ ID NO:38

Human MT1-MMP intron 1s

20 bp

5'CTCGGCTCGGCCCAAAGCAG3'

SEQ ID NO:39

Human MT1-MMP intron 1a

20 bp

5'GTAGGTCCCCGGGAGGCAGG 3'

SEQ ID NO:40

Human MT1-MMP intron 2s

20 bp

5'GTTTTACGGCTTGCAAGTAAC 3'

SEQ ID NO:41

Human MT1-MMP intron 2a

20 bp

5'CCAAACTTGTCTGGAACACC 3'

SEQ ID NO:42

Human MT1-MMP intron 3s

20 bp

5'CCAGGGTCTCAAATGGCAAC 3'

SEQ ID NO:43

Human MT1-MMP intron 3a

20 bp

5'ATGTGGCATACTCGCCCACC 3'

SEQ ID NO:44

Human MT1-MMP intron 4s

20 bp

5'CTCTGCCGAGCCTTGGACTG 3'

SEQ ID NO:45

Human MT1-MMP intron 4a

20 bp

5'GCATGGCCCAGCTCGTGCAC 3'

SEQ ID NO:46

Human MT1-MMP intron 5s

20 bp

5'TGCCCCGATGATGACCGCCGG 3'

SEQ ID NO:47

Human MT1-MMP intron 5a

20 bp

5'GGGTTGAGGGGGGCATCTTGG 3'

SEQ ID NO:48

Human MT1-MMP intron 6s

20 bp

5'CACCGTGGCCATGCTCCGAG 3'

SEQ ID NO:49

Human MT1-MMP intron 6a

20 bp

5'CCATCACTTGGTTATTCCTC 3'

SEQ ID NO:50

Human MT1-MMP intron 7s

20 bp

5'CCTACGAGAGGAAGGATGGC 3'

SEQ ID NO:51

Human MT1-MMP intron 7a

20 bp

5'GGTTCCAGGGACGCCTCATC 3'

SEQ ID NO:52

Human MT1-MMP intron 8s

20 bp

5'GGATGCCCAATGGAAAGACC 3'

SEQ ID NO:53

Human MT1-MMP intron 8a

20 bp

5'CGCTATCCACTGCCCTGAGC 3'

SEQ ID NO:54

Human MT1-MMP intron 9s

20 bp

5'GGGATCCCTGAGTCTCCCAG 3'

SEQ ID NO:55

Human MT1-MMP intron 9a

20 bp

5'TGTTGAATTTCCAGTATTTG 3'

SEQ ID NO:56

Human MT1-MMP Promoter 5s-1 (-480 )

20 bp

5'-TATTAGTAAACTGGCCCTTC-3'

SEQ ID NO:57

Human MT1-MMP Promoter 3a

20 bp

5'-ATCTTTCTTCTGCTTAGTCG-3'

SEQ ID NO:58

Human MT1-MMP Promoter 5s-2 (-790)

20 bp

5'-TAGAGGTGGAATAAACCCC-3'

SEQ ID NO: 59

Human MT1-MMP exon 5 PCR product  
285 bp

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1 GGGAGGCTGA GGGAAGGGAC TCAGGCTGCT ATCGTCACTG TCCCCATCCTT
51 CCAGGAAATG ACATCTTCCT GGTGGCTGTG CACGAGCTGG GCCATGCCCT
101 GGGGCTCGAG CATTCCAGTG ACCCTCGGC CATCATGGCA CCCTTTTACC
151 AGTGGATGGA CACGGAGAAT TTTGTGCTGC CCGATGATGA CCGCCGGGGC
201 ATCCAGCAAC TTTATGGCGA GTAGTCTACA CCCACGCCTG CTCCTCCTC
251 TGCTGCTTGT TCCCTCCTGG TCTACGCATT TCCCC

```

SEQ ID NO: 60

Human MT1-MMP exon 5 PCR product with P259P polymorphism  
285 bp

```

1 GGGAGGCTGA GGGAAGGGAC TCAGGCTGCT ATCGTCACTG TCCCCATCCTT
51 CCAGGAAATG ACATCTTCCT GGTGGCTGTG CACGAGCTGG GCCATGCCCT
101 GGGGCTCGAG CATTCCAGTG ACCCTCGGC CATCATGGCA CCGTTTACC
151 AGTGGATGGA CACGGAGAAT TTTGTGCTGC CCGATGATGA CCGCCGGGGC
201 ATCCAGCAAC TTTATGGCGA GTAGTCTACA CCCACGCCTG CTCCTCCTC
251 TGCTGCTTGT TCCCTCCTGG TCTACGCATT TCCCC

```

SEQ ID NO: 61

Human MT1-MMP exon 5 PCR product with D273N polymorphism  
285 bp

```

1 GGGAGGCTGA GGGAAGGGAC TCAGGCTGCT ATCGTCACTG TCCCCATCCTT
51 CCAGGAAATG ACATCTTCCT GGTGGCTGTG CACGAGCTGG GCCATGCCCT
101 GGGGCTCGAG CATTCCAGTG ACCCTCGGC CATCATGGCA CCGTTTACC
151 AGTGGATGGA CACGGAGAAT TTTGTGCTGC CCAATGATGA CCGCCGGGGC
201 ATCCAGCAAC TTTATGGCGA GTAGTCTACA CCCACGCCTG CTCCTCCTC
251 TGCTGCTTGT TCCCTCCTGG TCTACGCATT TCCCC

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SEQ ID NO: 62

Human ABCR cDNA (GenBank# NM\_000350)  
7318 bp

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1 ctggctctta acggcggtta tgcctttgc tgtctgaggg gcctcagctc tgaccaatct
61 ggtcttcgtg tggcattag catgggttc gtgagacaga tacagctttt gctctggaag
121 aactggaccc tgcggaaaag gcaaaagatt cgtttgtgg tggaactcgt gtggccttta
181 tctttatttc tggcttgat ctggttaagg aatgccaacc cgctctacag ccatcatgaa
241 tgccatttcc ccaacaaggc gatgccctca gcaggaatgc tgccgtggct ccaggggatc
301 ttctgcaatg tgaacaatcc ctgtttcaa agccccccc caggagaatc tcttgaatt
361 gtgtcaaaact ataacaactc catcttgga agggatatc gagattttca agaactctc
421 atgaatgcac cagagagcca gcaccttggc cgtatttga cagagctaca catctgtcc
481 caattcatgg acacctccg gactacccg gagagaattg caggaagagg aatacgaata
541 agggatatct tgaagatga agaaactg acactatttc tcattaaaaa catcggcctg
601 tctgactcag tggctacct tctgatcaac tctcaagtc gtccagagca gttcgtcat

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661 ggagtcctcg accctggcgt gaaggacatc gcctgcagcg aggccctcct ggagcgttc  
 721 atcatcttca gccagagacg cggggcaaag acgggtcgct atgccctgtg ctccctctcc  
 781 cagggcaccc tacagtggat agaagacact ctgtatgcca acgtggactt cttcaagctc  
 841 ttccgtgtgc tccccacact cctagacagc cgttctcaag gtatcaatct gagatcttgg  
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 1201 atccagagcc tggagtcaaa tcctttaacc aaaatcgctt ggagggcggc aaagcctttg  
 1261 ctgatgggaa aaatcctgta cactcctgat tcacctgcag cacgaaggat actgaagaat  
 1321 gccaaactca cttttgaaga actggaacac gttagggaagt tggtaaagc ctgggaagaa  
 1381 gtaggggccc agatctggta ctctttgac aacagcacac agatgaacat gatcagagat  
 1441 accctgggga acccaacagt aaaagacttt ttgaataggc agcttgggtga agaaggtatt  
 1501 actgctgaag ccactctaaa ctctctac aaggggcctc gggaaagcca ggctgacgac  
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 1621 aatcaatacc tggagtgtt ggtcctggat aagtttgaag gctacaatga tgaaactcag  
 1681 ctacccaac gtgccctctc tctactggag gaaaacatgt tctgggccgg agtggtattc  
 1741 cctgacatgt atccctggac cagctctcta ccacccacg tgaagtataa gatccgaatg  
 1801 gacatagacg tgggtggagaa aaccaataag attaaagaca ggtattggga ttctggtccc  
 1861 agagctgac ccgtggaaga ttccgggtac atctggggcg ggtttgcta tctgcaggac  
 1921 atggttgaac aggggatcac aaggagccag gtgcaggcg aggtccagt tggatctac  
 1981 ctccagcaga tgcctaccc ctgctcgtg gacgattct tcatgatcat cctgaaccgc  
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 2161 gcagtgtatt ggtgtacctg gttcctggac agcttctcca tcatgtcgat gagcatcttc  
 2221 ctctgacga tattcatcat gcatggaaga atcctacatt acagcgaccc attcatctc  
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 2341 ttcttctcca aggccagtct ggcagcagcc tgtagtggg tcatctattt caccctctac  
 2401 ctgccacaca tcctgtgctt cgctggcag gaccgcatga ccgctgagct gaagaaggct  
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 2521 gagcaaggcc tggggctgca gtggagcaac atcggaaga gtcccacgga aggggacgaa  
 2581 ttacgttcc tctgtccat gcagatgatg ctcttgatg ctgcgtgcta tggcttactc  
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 2821 cagactcct tctttgaacg tgagcatcca ggtgggttc ctggggtatg cgtgaagaat  
 2881 ctggtaaaga ttttgagcc ctgtggccgg ccagctgtgg accgtctgaa catcaccttc  
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 3001 tccatcctga cgggtctgtt gccaccaacc tctgggactg tctcgttgg gggaagggac  
 3061 attgaaacca gcctggatgc agtccggcag agccttggca tgtgtccaca gcacaacatc  
 3121 ctgttccacc acctcaggtt ggctgagcac atgctgttct atgccagct gaaaggaaag  
 3181 tcccaggagg agggccagct ggagatggaa gccatgttgg aggacacagg cctccaccac  
 3241 aagcggaatg aagaggctca ggacctatca ggtggcatgc agagaaagct gtcggttgc  
 3301 attgcctttg tgggagatgc caaggtgtg attctggacg aaccacctc tggggtggac  
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3421 atcatgccc ctcaccacat ggacgaggcc gaccaccaag gggaccgcat tgccatcatt  
 3481 gcccaggga ggctctactg ctccaggacc ccactctcc tgaagaactg ctttggcaca  
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 3601 gaggggacct gcagctgctc gtctaagggt ttctccacca cgtgtccagc ccacgtcgat  
 3661 gacctaac cagaacaagt cctggatggg gatgtaatg agctgatgga ttagttctc  
 3721 caccatgttc cagaggcaaa gctggtggag tgcatgttc aagaacttat ctctctctt  
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 3961 agagaaaacg tcaacccccg acaccctgc ttgggtccca gagagaaggc tggacagaca  
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 4201 ctggcgacga tctgtctccc ggctacctt gtgttttg ctctgatgct ttctattgtt  
 4261 atcttctct ttggcgaata ccccgcttg acccttcacc cctggatata tgggcagcag  
 4321 tacaccttct tcagcatgga tgaaccaggc agtgagcagt tcacggtact tgcagacgtc  
 4381 ctctgaata agccaggctt tggcaaccgc tgcctgaagg aagggtggct tccggagtag  
 4441 ccctgtggca actcaacacc ctggaagact cttctgtgt ccccaaacat caccagctg  
 4501 ttccagaagc agaatggac acaggtaac ccttcacat cctgcagggt cagcaccagg  
 4561 gagaagctca ccatgtgcc agagtgcacc gaggtgccc ggggcctccc gccccccag  
 4621 agaacacagc gcagcacgga aattctaca gacctgacgg acaggaacat ctccgacttc  
 4681 ttgtaaaaa cgtatcctgc tcttataaga agcagctta agagcaaatt ctgggtcaat  
 4741 gaacagaggt atggaggaat ttccattgga gaaagctcc cagtcgtccc catcacgggg  
 4801 gaagcacttg ttgggtttt aagcgacct ggccggatca tgaatgtgag cggggggcct  
 4861 atcactagag aggcctctaa agaaatacct gatttctta aacatctaga aactgaagac  
 4921 aacattaagg tgtggtttaa taacaaaggc tggcatgccc tggtcagctt tctaatgtg  
 4981 gcccacaacg ccatcttac ggccagcctg cctaaggaca ggagccccga ggagtatgga  
 5041 ataccgtca ttagccaacc cctgaacctg accaaggagc agctctcaga gattacagt  
 5101 ctgaccactt cagtggatgc tgtggtgcc atctgcgtga ttttccat gtccttcgtc  
 5161 ccagccagct ttgtcttta ttgatccag gagcgggtga acaaatcaa gcacctccag  
 5221 ttatcagtg gactgagccc caccacctac tgggtgacca acttctctg ggacatcatg  
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 5341 tacattctc cagaaaacct tctgacctt gtggcactgc tctgctgta tggatggcg  
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 5521 gaattattg ataataaccg gacgtgctc aggttcaacg ccgtgctgag gaagctgctc  
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 5641 gtgacagatg tctatcccg gtttggtag gagcactctg caaatccgtt cactgggac  
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 5761 ctgctgtcc agcgcactt ctctctcc caatggattg ccgagcccac taaggagccc  
 5821 attgtgatg aagatgatga tgtggctgaa gaaagacaaa gaattattac tgggtgaaat  
 5881 aaaactgaca tcttaaggct acatgaacta accaagattt atctgggcac ctccagccca  
 5941 gcagtggaca ggctgtgtg cggagttgc cctggagagt gctttggcct cctgggagt  
 6001 aatggtgccg gcaaaacaac cacattcaag atgtcactg gggacaccac agtgacctca  
 6061 ggggatgcca ccgtagcagg caagagtatt ttaaccaata ttctgaagt ccatcaaat  
 6121 atgggctact gtcctcagtt tgatgcaat gatgagctgc tcacaggacg agaacatctt

6181 tacctttatg cccggcttcg aggtgtacca gcagaagaaa tcgaaaagggt tgcaaaactgg  
6241 agtattaaga gcctgggcct gactgtctac gccgactgcc tggctggcac gtacagtggg  
6301 ggcaacaagc ggaaactctc cacagccatc gcactcattg gctgcccacc gctggtgctg  
6361 ctggatgagc ccaccacagg gatggacccc caggcacgcc gcatgctgtg gaacgtcatc  
6421 gtgagcatca tcagaaaagg gagggctgtg gtctcacat cccacagcat ggaagaatgt  
6481 gaggcactgt gtacccggct ggccatcatg gtaaagggcg ccttcgatg tatgggcacc  
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6601 ccgaaggacg acctgcttcc tgacctgaac cctgtggagc agttcttcca ggggaacttc  
6661 ccaggcagtg tgcagaggga gaggcactac aacatgctcc agttccaggt ctctcctcc  
6721 tccctggcga ggatcttcca gctcctctc tcccacaagg acagcctgct catcgaggag  
6781 tactcagtca cacagaccac actggaccag gtgtttgtaa attttgctaa acagcagact  
6841 gaaagtcatg acctccctct gcacctcga gctgtggag ccagtcgaca agcccaggac  
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6961 ggagcctgtg cccatatggt catccaatg gactggccca gcgtaatga cccactgca  
7021 gcagaaaaca aacacacgag gagcatgcag cgaattcaga aagaggtctt tcagaaggaa  
7081 accgaaactg acttgctcac ctggaacacc tgatggtgaa accaaacaaa taaaaatcc  
7141 ttctccagac ccagaaacta gaaaccccg gccatccac tagcagctt ggctccata  
7201 ttgtctcat ttcaagcaga tctgctttc tgcattttg tctgtgtgc tgcgttgtg  
7261 gtgatttca tggaaaaata aatgcaaat gcatcatca caaaaaaaaa aaaaaaaaa

SEQ ID NO: 63

Human apolipoprotein E cDNA (GenBank# NM\_000041)

1156 bp

1 cgcagcggag gtgaaggacg tccttcccca ggagccgact ggccaatcac aggcaggaag  
61 atgaaggttc tgtgggctgc gttgctgtgc acattcctgg caggatgcca ggccaaggtg  
121 gagcaagcgg tggagacaga gccggagccc gagctgcgcc agcagaccga gtggcagagc  
181 ggccagcgtg gggaactggc actgggtcgc ttttgggatt acctgcgctg ggtgcagaca  
241 ctgtctgagc aggtgcagga ggagctgctc agtcccagg tcaccagga actgaggcgc  
301 ctgatggacg agaccatgaa ggagttgaag gcctacaaat cggaaactgga ggaacaactg  
361 acccgggtgg cggaggagac gcgggcacgg ctgtccaagg agctgcaggc ggcgagggc  
421 cggctgggcg cggacatgga ggacgtgtgc ggccgctgg tcagttaccg cggcgaggtg  
481 caggccatgc tcggccagag caccgaggag ctgcgggtgc gcctgcctc ccactgcgc  
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601 caggccgggg cccgcgaggg cgcgagcgc ggctcagcg ccatccgcga gcgcctgggg  
661 cccctggttg aacagggccg cgtgcgggcc gccactgtgg gctccctggc cggccagccg  
721 ctacaggagc gggcccaggc ctggggcgag cggctgcgcg cgcggatgga ggagatgggc  
781 agccggaccc gcgaccgctt ggacgaggtg aaggagcagg tggcggaggt gcgcgccaag  
841 ctggaggagc agggccagca gatagcctg caggccgagg cctccaggc ccgcctcaag  
901 agctggttcg agcccctggt ggaagacatg cagcgccagt gggccgggct ggtggagaag  
961 gtgcaggctg ccgtgggcac cagcgccgcc cctgtgcca gcgacaatca ctgaacgccg  
1021 aagcctgcag ccatgcgacc ccacgccacc ccgtgcctcc tgcctccgcg cagcctgcag  
1081 cgggagaccc tgtcccgccc ccagccgtcc tctgggggtg gaccctagt taataaagat  
1141 tcaccaagtt tcacgc

SEQ ID NO: 64

Human C-C chemokine receptor-2 (Ccr-2) cDNA (GenBank# NM\_000647)  
2273 bp

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1 caggactgcc tgagacaagc cacaagctga acagagaaag tggattgaac aaggacgcat
61 ttccccagta catccacaac atgctgtcca catctcgttc tcggtttatc agaaatacca
121 acgagagcgg tgaagaagtc accacctttt ttgattatga ttacgggtgct ccctgtcata
181 aatttgacgt gaagcaaatt ggggcccaac tcctgcctcc gctctactcg ctgggtgtca
241 tctttgggtt tgtgggcaac atgctggctg tcctcatctt aataaactgc aaaaagctga
301 agtgcttgac tgacatttac ctgctcaacc tggccatctc tgatctgctt ttcttatta
361 ctctccatt gtgggtcac tctgtgcaa atgagtgggt ctttgggaat gcaatgtgca
421 aattattcac agggctgtat cacatcggtt attttggcgg aatcttcttc atcatcctcc
481 tgacaatcga tagatactg gctattgtcc atgctgtgtt tgcttataaa gccaggacgg
541 tcacctttgg ggtggtgaca agtgtgatca cctggttggg ggctgtgttt gcttctgtcc
601 caggaatcat ctttactaaa tgccagaaag aagattctgt ttatgtctgt ggcccttatt
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721 cgctgctcat catggtcatc tgctactcgg gaatcctgaa aacctgtctt cggtgtcgaa
781 acgagaagaa gaggcatagg gcagtgagag tcactctcac catcatgatt gtttacttcc
841 tctctggac tcctataaac attgtcattc tcctgaacac ctccaggaa ttctcggcc
901 tgagtaactg tgaagcacc agtcaactgg accaagccac gcaggtgaca gagactcttg
961 ggatgactca ctgctgcac aatcccatca tctatgcctt cgttggggag aagttcagaa
1021 gcctttttca catagctctt ggctgtagga ttgccccact caaaaacca gtgtgtggag
1081 gtccaggagt gagaccagga aagaatgtga aagtgactac acaaggactc ctcatggtc
1141 gtggaaaagg aaagtcaatt ggagagccc ctgaagccag tcttcaggac aaagaaggag
1201 cctagagaca gaaatgacag atctctgctt tggaaatcac acgtctggct tcacagatgt
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1321 gactccagct gggttgaaa acagtatttt ccaaactacc ttccagtcc tcattttga
1381 atacaggcat agagttcaga cttttttaa atagtaaaaa taaaattaaa gctgaaaact
1441 gcaacttgta aatgtggtaa agagttagtt tgagttgcta tcatgtcaaa cgtgaaaatg
1501 ctgtattagt cacagagata attctagctt tgagcttaag aattttgagc aggtgggatg
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1621 tctgtgggca cattagccta tgtcatgca gcatctaagt aatgatgtcg ttgaatcac
1681 agtatacgtc ccctcgtgt catctcagct ggatcctcat tctctcaggc ttgctgcaa
1741 aagccttttg tgtttgttt tgtatcatta tgaagtcag cgtttaatca cattcagtg
1801 ttacgtgct tcgcagatgt cctgatgct catattgttc cctaatttgc cagtgggaac
1861 tcctaaatca aattggcttc taatcaaagc ttttaaacc tatttgtaaa gaatggaagg
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1981 tctatgttg ccagtggtt ttctgatctg atgcaagcaa gaaacactgg gcttctagaa
2041 ccaggcaact tgggaactag actcccaagc tggactatgg ctctacttcc aggccacatg
2101 gctaaagaag gtttcagaaa gaagtgggga cagagcagaa cttcacctt catatatttg
2161 tatgatccta atgaatgcat aaaatgttaa gttgatgtg atgaaatga aatactgttt
2221 ttaacaacta tgatttgaa aataaatcaa tgctataact atgttgataa aag

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SEQ ID NO: 65

Human cystatin C cDNA (GenBank# NM\_000099)

818 bp

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1 cgcagcgggt cctctctatc tagctccagc ctctgcctg cgccccactc cccgcgtccc
61 gcgtcctagc cgacatggc cgggccccctg cgcgccccgc tgctcctgct ggccatcctg
121 gccgtggccc tggccgtgag ccccgcgggc ggctccagtc ccggcaagcc gccgcgcttg
181 gtgggaggcc ccatggacgc cagcgtggag gaggagggtg tgcggcgtgc actggacttt
241 gccgtcggcg agtacaacaa agccagcaac gacatgtacc acagccgcgc gctgcagggtg
301 gtgcgcgccc gcaagcagat cgtagctggg gtgaactact tcttgacgt ggagctgggc
361 cgaaccacgt gtaccaagac ccagcccaac ttggacaact gcccttcca tgaccagcca
421 catctgaaaa ggaaagcatt ctgctcttc cagatctacg ctgtgccttg gcagggcaca
481 atgacctgt cgaatccac ctgtcaggac gcctaggggt ctgtaccggg ctggcctgtg
541 cctatcact cttatgcaca cctccaccc cctgtattcc caccctgga ctggtggccc
601 ctgccttggg gaaggtctcc ccatgtgcct gcaccaggag acagacagag aaggcagcag
661 gcggcctttg ttgctagca aggggctctg cctccctcc ttccttctg ctctcatag
721 ccccggtgtg cgggtgcatc accccacct cctgcaataa aatagtagca tcggcaaaaa
781 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa

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SEQ ID NO: 66

Human hemicentin/FIBL-6 cDNA (GenBank# NM\_031935)

18209 bp

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1 gaagccgcat ccagacaaaa gctgccgcat cctgccctg cccaaccct ggagggattc
61 gagtttggtg cttgtcccg tctgattctc agcgccaaac ttttgctag ttcagagatt
121 ccaagagtct gatgagttac tctgagagga aaccctctgc ctgttggtga ggaggactga
181 gcacagtgtc taggcgtga ggggggaaaa gagggggaaa aaaaagaaaa tgatttctg
241 ggaagttgtc catacagtat tctgtttgc tcttctttat tcttcctag ctcaagatgc
301 gagccccag tcagagatca gagctgagga aattcccgag ggggcctcca cgttggttt
361 tgtgttgat gtgactggtt ctatgtatga tgattagt caggtgattg aaggggctc
421 caaaatttg gagacgtctt tgaaaagacc taaaagacct ctttcaact ttgcgttggt
481 gcctttccat gatccagaaa ttggccagc gacaattacc acagatccca agaaatttca
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601 agctataaaa attgccttg aaatttctc tctggttct tcatctatg tttcactga
661 tgctcggtcc aaagattacc ggctcaccca tgagtgctg caacttatec aacagaaaca
721 gtcacaagtc gtattgttc tgactggaga ttgtgatgac aggaccata ttgatataa
781 agtctatgaa gaaattgcct ctacaagttc tggtaagtg ttcatctgg aaaaaaaca
841 agttaatgag gtattaaaat gggtagaaga agcagtacag gcctccaaag ttacctttt
901 atccacagat catttgaac aggctgtaaa tacttgagga attccttttg atccagcct
961 gaaagaggtc actgtgtctt tgagtgggcc ttctccaatg attgaaattc gcaatcctt
1021 agggaaagctg ataaaaaagg gatttggcct gcatgagcta ttaaataatc ataactctgc
1081 caaagtagtg aatgtgaaag agccagaggc tggaatgtgg acagtgaaga cctcaagcag
1141 tggaaggcac tctgttcgca ttactggcct cagtactatt gattccgag ctggctttc
1201 tcgaaagccc accctggact tcaaaaaaac agtcagcaga ccagtgaag gaatacctac
1261 ctatgtactg ctcaatactt ctggaatttc cactccagct agaatagatc ttcttgaact
1321 tttagtate tcaggaagtt ctcttaagac tattcctgtt aaatattacc cacatcgaaa

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1381 accttatggc atatggaata ttctgactt tgtaccacca aatgaagctt tctttctcaa  
 1441 agtaacaggc tatgataaag atgattacct ctccagaga gtatcaagtg ttctctttc  
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 15241 cactgtaaag gattacacag aggactacat tcaaacaggt cctgggcagc tgtacgccta  
 15301 ctcaaccggg ctgttcacca ttgatggcat cagcatccca tacacatgga accacaccgt  
 15361 ttctatgat caggcacagg gaagaatgcc ttcttggtt gaaacacttc atgcacctc  
 15421 tgtggaatct gactataacc agatagaaga gacactgggt tttaaaatc atgcttcaat  
 15481 atccaaagga gatcgagta atcagtcccc ctccggggtt accttagact cagttggacc  
 15541 ttttgtgct gatgaggatg aatgtgcagc agggaaatccc tgctccata gctgccacia  
 15601 tgccatgggg acttactact gctcctgcc taaaggcctc accatagctg cagatggaag  
 15661 aacttgtcaa gatattgatg agtgtgctt gggtaggcat acctgccacg ctggtcagga  
 15721 ctgtgacaat acgattggat ctatcgctg tgtggtccgt tgtggaagt gctttcgaag  
 15781 aacctctgat gggctgagtt gtaagatat taatgaatgt caagaatcca gccctgtca  
 15841 ccagcgctgt ttcaatgcca taggaagttt ccattgtgga tgtgaacctg ggtatcagct  
 15901 caaaggcaga aatgcatgg atgtgaacga gtgtagacaa aatgtatgca gaccagatca  
 15961 gcactgtaag aacaccgtg gtggctataa gtgcattgat cttgtccaa atggaatgac  
 16021 caaggcagaa aatggaacct gtattgatat tgatgaatgt aaagatggga cccatcagtg  
 16081 cagatataac cagatatgtg agaatacaag aggcagctat cgttgtgtat gccaaagg  
 16141 ttatcggtct caaggagtg gaagaccctg catggacatt aatgaatgtg aacaagtgcc  
 16201 taaacctgtg gcacatcagt gctccaacac cccggcagc ttcaagtga tctgtccacc  
 16261 aggacaacat ttattagggg acgggaaatc ttgcgtgga ttggagaggc tgccaaatta  
 16321 tggcactcaa tacagtagct ataacctgc acggttctcc cctgtgagaa acaactatca  
 16381 acctcaacag cattacagac agtactaca tctctacagc tctactcag agtatagaaa  
 16441 cagcagaaca tctctctcca ggactagaag gactattagg aaaacttgcc ctgaaggctc  
 16501 tgaggcaagc catgacacat gtgtagatat tgatgaatgt gaaaatacag atgcctgcca  
 16561 gcatgagtgt aagaatacct ttggaagta tcagtgcac tgcccacctg gctatcaact  
 16621 cacacacaat ggaaagacat gccaagatat cgatgaatgt ctggagcaga atgtgcactg  
 16681 tggaccaat cgcatgtgct tcaacatgag aggaagctac cagtgcacg atacaccctg  
 16741 tccaccaac taccaacggg atcctgttc agggttctgc ctcaagaact gtccaccaa  
 16801 tgatttgaa tgtgcctga gcccatatgc ctggaatac aaactcgtc cctcccatt  
 16861 tggaatagcc accaatcaag attaatccg gctggttgca tacacacagg atggagtgtat  
 16921 gcatcccagg acaactttc tcatggtaga tgaggacag actgttctt ttgccttgag  
 16981 ggatgaaaac ctgaaaggag tgggtatatac aacacgacca ctacgagaag cagagacct  
 17041 ccgcatgagg gtccgagcct cactctacag tgccaatggg accattgaat atcagaccac  
 17101 attcatagtt tatatagctg tgccgccta tccatactaa ggaactctc aaagcctatt  
 17161 ccacatattt aaaccgcat aatcatggca atcaagcccc ctccagatt actgtctt  
 17221 gaacagttgc aatcttgga gcttgaaaat ggtgctacac tctgtttgt gtgccttct  
 17281 tggctactct gaggtattt catgatccca ccatggcat atctgaagt atggtctaga  
 17341 aaagtccctt attattttat ttattact ggagcagta ctcccaaag attattctga  
 17401 acatctaaca ggacatatca gtgatggtt acagtgtgt agtacctaag atcatttcc  
 17461 tgaagccaa accaaacaac gaaaacaag aacaactaat tcagaatcaa atagagttt  
 17521 tgagcatttg actattttta gaatcataaa attagtact aagtatttg atcaaagctt  
 17581 ataaaataac ttacggagat tttgtaagt attgatacat tataatagga ctgcctatt  
 17641 ttcatttta agaagaaaa caccactcat ttataaaa atagtacagc tactataagg  
 17701 ctgtttgat cccaaatggg gcttacttg attgaacatt cagaacaagg atattattt  
 17761 cagtgtttt gtgagatcag ctgaaccact tatgataata ataataaaa agactgctt  
 17821 gccctcagc cagtgtaca tggcatggaa ctttaaaa tttaataata actttcatcc  
 17881 agttagcttc ataacttta cgttcagaa tttgtttat ttctgtca atgaaagca

17941 tttttaaaga taccagtggg acaggtttgg tttttaaaa atctcatgtg ttcaaattaa  
 18001 cataaatatt acacgtcaat aactgtaca tgggtgtaat agactctaag caattgccaa  
 18061 gatgtattct attttatga agtgtatata tattacctta gtgtgcattt tctatataat  
 18121 atcttgatgg actcttttat aaaattattt tataaaaaac aatgttacac taaaatcagc  
 18181 ctaaataaat ttccacaact tttttcat

SEQ ID NO: 67

Human manganese superoxide dismutase 2 cDNA (GenBank# NM\_000636)  
 1026 bp

1 cagcatgttg agccgggcag tgtgcggcac cagcaggcag ctgcctccgg ttttggggta  
 61 tctgggctcc aggcagaagc acagcctccc cgacctgccc tacgactacg gcgcccctgga  
 121 acctcacatc aacgcgcaga tcatgcagct gcaccacagc aagcaccacg cggcctacgt  
 181 gaacaacctg aacgtcaccg aggagaagta ccaggaggcg ttggccaagg gagatgttac  
 241 agcccagata gctcttcagc ctgcactgaa gtccaatggt ggtggtcata tcaatcatag  
 301 cattttctgg acaaacctca gccctaacgg tgggtggagaa ccaaaggagg agttgctgga  
 361 agccatcaaa ctggactttg gttcctttga caagttaag gagaagctga cggtgcac  
 421 tgttggtgtc caaggctcag gttggggttg gcttggttc aataaggaaac ggggacactt  
 481 acaaatgct gcttgctcaa atcaggatcc actgcaagga acaacaggcc ttattccact  
 541 gctggggatt gatgtgtggg agcacgctta ctacctcag tataaaaatg tcaggcctga  
 601 ttatctaaaa gctatttga atgtaataca ctgggagaat gtaactgaaa gatacatgac  
 661 ttgcaaaaag taaaccacga tcgttatgct gagtatgta agctctttat gactgtttt  
 721 gtagtggtat agagtactgc agaatacagt aagctgctct attgtagcat ttcttgatgt  
 781 tgcttagtca ctatttcat aaacaactta atgttctgaa taatttctta ctaaacattt  
 841 tgttattggg caagtgttg aaaatagtaa atgctttgtg tgattgaatc tgattggaca  
 901 tttcttcag agagctaaat tacaattgtc attataaaa ccatcaaaaa tattccatcc  
 961 atatacttg gggacttgta gggatgcctt tctagtccta ttctattgca gttatagaaa  
 1021 atctag

SEQ ID NO: 68

Human C-C chemokine ligand 2 (Ccl-2)/monocyte chemoattractant protein 1 cDNA  
 (GenBank# NM\_002982)  
 757 bp

1 ggaaccgaga ggctgagact aaccagaaa catccaattc taaaactgaa gctcgcactc  
 61 tcgcctccag catgaaagtc tctgccgccc ttctgtgcct gctgctcata gcagccacct  
 121 tcattcccca agggctcgtc cagccagatg caatcaatgc ccagtcacc tgctgttata  
 181 acttcaccaa taggaagatc tcagtgcaga ggctcgcgag ctatagaaga atcaccagca  
 241 gcaagtgtcc caaagaagct gtgatcttca agaccattgt ggccaaggag atctgtgctg  
 301 accccaagca gaagtgggtt caggattcca tggaccacct ggacaagcaa acccaaactc  
 361 cgaagacttg aactcact ccacaacca agaactcga gctaacttat tttcccctag  
 421 ctttcccag acaccctgtt ttattttatt ataataaatt ttgtttgtg atgtgaaaca  
 481 ttatgcctta agtaatgta attctattt aagtattga tgtttaagt ttatcttca  
 541 tggtagtagt gttttttaga tacagagact tggggaaatt gcttttcctc ttgaaccaca  
 601 gttctacccc tgggatgtt tgagggtctt tgcaagaatc attaatataa agaattttt  
 661 ttaacattcc aatgcattgc taaaatatta ttgtggaaat gaatttttg taactattac

721 accaaataaa tatatttttg tacaaaaaaa aaaaaaa

SEQ ID NO: 69

Human paraoxonase 1 cDNA (GenBank# NM\_000446)

2395 bp

1 agagcctcct agcccgtcgg tgtctgcgcc catcgatccc ttgtctatc cccgaccatg  
 61 gcgaagctga ttgcgtcac cctcttgggg atgggactgg cactcttcag gaaccaccag  
 121 tcttctacc aaacacgact taatgtcttc cgagaggtag aaccgtaga acttcctaac  
 181 tgtaatttag ttaaaggaat cgaaactggc tctgaagact tggagatact gcctaattga  
 241 ctggcttca ttagctctgg attaaagtat cctggaataa agagcttcaa cccaacagt  
 301 cctggaaaaa tacttctgat ggacctaagt gaagaagatc caacagtgtt ggaattgggg  
 361 atcactggaa gtaatttga tgtatctca tttaaccctc atgggattag cacattcaca  
 421 gatgaagata atgcatgta cctcctgggt gtgaaccatc cagatgccaa gtccacagt  
 481 gagttgttta aatttcaaga agaagaaaaa tcgcttttgc atctaaaaac catcagacat  
 541 aaactctgc ctaatttga tgatattgtt gctgtgggac ctgagcactt ttatggcaca  
 601 aatgatcact atttcttga cccctactta caatcctggg agatgtattt gggtttagcg  
 661 tggctgatg ttgtctacta tagtccaagt gaagttcgag tgggtggcaga aggattgat  
 721 ttgctaagt gaatcaacat ttacccgat ggcaagtatg tctatatagc tgagttgctg  
 781 gctcataaga ttcatgtgta tgaaaagcat gctaattgga cttaactcc attgaagtcc  
 841 ctgacttta ataccctcg ggataacata tctgtggatc ctgagacagg agaccttgg  
 901 gttggatgcc atcccaatgg catgaaaatc ttctctatg actcagagaa tctcctgca  
 961 tcagaggtgc ttgaatcca gaacattcta acagaagaac ctaaagtac acaggtttat  
 1021 gcagaaaatg gcacagtgtt gcaaggcagt acagttgcct ctgtgtacaa agggaaactg  
 1081 ctgattggca cagtgttca caaagctctt tactgtgagc ttaacagac cgatttgcac  
 1141 ccatgccata gaaactgagg ccattatttc aaccgcttgc catattccga ggaccagtg  
 1201 ttcttagctg aacaatgaat gctgacccta aatgtggaca tcatgaagca tcaaagcact  
 1261 gttaactgg gagtgatatg atgtgtaggg ctttttttg agaatacact atcaaatcag  
 1321 tcttgaata ctgaaaacc tcattacca taaaaatcct tctactaaa atggataaat  
 1381 cagttatgtc aattgtcaga tattaaataa cagtgtgtga ccccaaaagt acttacccta  
 1441 aaacatgtgt tgcctgaaag cacatgtgtg tatcgtgcc ttgccatgtc ttgttcagaa  
 1501 gacacagggg agcaggggta gctcacgtgt cttagaact ccagtactca cccagggact  
 1561 ccagttcaca ggccagaaaa catatgcatt atgaagtcc cctctactcc atgcacatag  
 1621 taagtctgac tatggcagtc agacttactt actccattt tcccttcgat atatgacttt  
 1681 ttctcagtaa atattaacct gaactattcc aactccccct gtactcttgc ttttcaatt  
 1741 ctctgttgc aatgacacat aggaaaatct taaaattctt gggagtgttg tcacacctga  
 1801 aaattatgag tctctatgat ctggcaciaa attgtacatt tgagtgtctt tgacttggtt  
 1861 aaagggaagt ttgtcacttc gatgactgga tacagaatga atcccataat tgacatgggc  
 1921 gacagtaaaa gtgtcccaa agactacact gttgtttagg tgggtgtagt gctggtgggt  
 1981 tttgtttaa tattaaact tctgtgtg gaggtgaaa agaaaaaaaa taatagaaag  
 2041 gtaacaaac aaataaatag aaaagatcaa caacccttt ggctatctac tgagacatga  
 2101 ctaggaagaa aacatgactt tatcattttg ttatagaagc tgatatataa ggttacacat  
 2161 ttcatattat ttgttttct gatttgaagg tataaccttc atgatgaatt acttcttcag  
 2221 ggtgttaagg cagtgttctt agaaacaaat tttttcttg cttttgtttt gttttgaga  
 2281 ccgaatctca ctctgttccc caggctggag tgcagtgggt cgatcttggc tcaactgcaac  
 2341 ttctacctcc gaggttcaag agattcttgt gcctcagcct cccggatagc tgccg

SEQ ID NO: 70

Human unknown protein PHG-1 hypothetical peptide 1 (GenBank # AL832747)

74 aa

MAFLVHSQPVILGFTVLLSYILRYQLLFFKFVFIKFDKKPALATHHNK  
SHFKIVAQTPRKKRKEKLEQQQQKN

SEQ ID NO: 71

Human unknown protein PHG-1 hypothetical peptide 2 (GenBank # AL832747)

55 aa

MTLLVFTSHVQCPNRQCKKYPVWFNRKSVYVSLFETSFTLSGSLSSM  
KSARNIGW

SEQ ID NO: 72

Human unknown protein PHG-1 hypothetical peptide 3 (GenBank # AL832747)

52 aa

METNFEVLLPFDLGLEYELLYNSYSYLANAQFSITSLMAFTRKAVLEA  
IVIH

SEQ ID NO: 73

Human unknown protein PHG-1 hypothetical peptide 4 (GenBank # AL832747)

45 aa

MYFAMKLPLGLIISIPLLRNVQMILYSTTLVPLCMTVRFFFFLLF

SEQ ID NO: 74

Human unknown protein PHG-1 hypothetical peptide 5 (GenBank # AL832747)

43 aa

MDRENQISSYNCLANGISGSFSASHFRLHSLTLLHFKIPAFIF

SEQ ID NO: 75

Human unknown protein PHG-1 hypothetical peptide 6 (GenBank # AL832747)

37 aa

MCCFGYTHSFFFNRIYCLVSLWTGTVD AHLKVKCHFF

SEQ ID NO: 76

Human unknown protein PHG-1 hypothetical peptide 7 (GenBank # AL832747)

35 aa

MFSVQTGNVKSILCGLTGNLFMSLYLKPVLLSVVL

SEQ ID NO: 77

Human unknown protein PHG-1 hypothetical peptide 8 (GenBank # AL832747)

34 aa

MIYFLKSNFNSSCLTEACQYMCCIFFAFVEKLHI

SEQ ID NO: 78

Human unknown protein PHG-1 hypothetical peptide 9 (GenBank # AL832747)

34 aa

MPRAIVFPFFASFSYPLFQLQMPKKMPTDTTLP

SEQ ID NO: 79

Human prostaglandin D2 synthase protein (GenBank# NM\_000954)

190 aa

MATHHTLWMGLALLGVLGDLQAAPEAQVSVQPNFQQDKFLGRWFSAGLASNS  
SWLREKKAALSMCKSVVAPATDGGLNLTSTFLRKNQCETRTMLLPAGSLGSYS  
YRSPHWGSTYSVSVVETDQYALLYSQGSKGPGEDFRMATLYSRTQTPRAEL  
KEKFTAFCKAQGFTEDTIVFLPQTDKCMTEQ

SEQ ID NO: 80

Human myelin basic protein (GenBank# M13577)

170 aa

MASQKRPSQRHGSKYLATASTMDHARHGFLPRHRDTGILDSIGRFGGDRGAPKR  
GSGKDSHHPARTAHYGSLPQKSHGRTQDENPVVHFFKNIVTPRTPPPSQGKGRG  
LSLSRFSWGAEGQRPFGYGGRASDYKSAHKGFKGVDAQGTLSKIFKLGGDRSR  
SGSPMARR

SEQ ID NO: 81

Human unknown protein PHG-4 peptide 1 (GenBank# AP006241)

38 aa

IRSAKLGFCCLNSALGPQINRCECSFFPLCEEAVTPQQ

SEQ ID NO: 82

Human unknown protein PHG-4 peptide 2 (GenBank# AP006241)

38 aa

LLGCNCFFTQGEKTTFTSVYLRTQCRVQAAKPQLSRSN

SEQ ID NO: 83

Human unknown protein PHG-4 peptide 3 (GenBank# AP006241)

37 aa

FIYKKIKLEIVLDFSSYCWGVTASSHRGKKLHSHRFI

SEQ ID NO: 84

Human unknown protein PHG-5 (GenBank# BC011973)

334 aa

MGASSSSALARLGLPARPWPRWLGVAAALGLAAVALGTVAWRRRAWPRRRRLQ  
QVGTVAKLWIYPVKSCKGVPVSEAECTAMGLRSGNLRDRFWLVIKEDGHMVTA  
RQEPRLVLISIIYENNCLIFRAPDMDQLVLPSKQPSSNKLHNCRIFGLDIKGRDCGN  
EAAKWFTNFLKTEAYRLVQFETNMKGRTSRKLLPTLDQNFQVAYPDYCPLLIMT  
DASLVDLNRMEKKMKMENFRPNIVVTGCDAFEEDTWDELLIGSVEVKKVMAC

PRCILTTVDPDTGVIDRKQPLDTLKSRYRLCDPSERELYKLSPLFGIYYSVEKIGSLR  
VGDPVYRMV

SEQ ID NO: 85

Human peanut-like 2/septin 4 protein (GenBank# NM\_080416)

459 aa

MIKRFLDITDDGELSKFVKDFSGNASCHPPEAKTWASRPQVPEPRPQAPDLYD  
DDLEFRPPSRPQSSDNQQYFCAPAPLSPSARPRSPWGKLDPYDSEDDKEYVGF  
ATLPNQVHRKSVKKGFDFTLMVAGESGLGKSTLVNSLFLTDLYRDRKLLGAEERI  
MQTVEITKHAVDIEEKGVRLRLTIVDTPGFGDAVNNTWCWPVAEYIDQQFEQY  
FRDEGLNRKNIQDNRVHCCLYFISPFHGHLRPLDVEFMKALHQRVNIVPILAKA  
DTLTPPEVDHKKRKIREEIEHFGIKIYQFPDCDSDEDEDFKLQDQALKESIPFAVIG  
SNTVVEARGRRVRGRLYPWGIVEVENPGHCDVFKLRTMLVRTHMQDLKDVTR  
THYENYRAQCIQSMTRLVVKERNRNKLTRESGTDFFIPAVPPGTDPETEKLIREK  
DEELRRMQEMLHKIQKQMKENY

SEQ ID NO: 86

Human coactosin-like 1 protein (GenBank# NM\_021149)

142 aa

MATKIDKEACRAAYNLVRDDGSAVIWVTFKYDGSTIVPGEQGAEYQHFIQQCTD  
DVRLFAFVRFTTGDAMSKRSKFALITWIGENVSGLQRAKTGTDKTLVKEVVQNF  
AKEFVISDRKELEEDFIKSELKKAGGANYDAQTE

SEQ ID NO: 87

Human clusterin protein (GenBank# BC019588)

449 aa

MMKTLLLFLVGLLLTWESGQVLGDQTVSDNELQEMSNQGSKYVNKEIQNAVNG  
VKQIKTLIEKTNEERKTLLSNLEEAKKKKEDALNETRESETKLKELPGVCNETMM  
ALWEECKPCLKQTCMKFYARVCRSGSLVGRQLEEFNLQSSPFYFWMNGDRID  
SLENDRQQTHMLDVMQDHFSRASSIIDELFQDRFFTREPQDITYHYLPFSLPHRR  
PHFFFFPKSRIVRSLMPFSPYEPLNFHAMFQPFLEMIHEAQQAMDIHFHSPAFQHPP  
TEFIREGDDDRTVCREIRHNSTGCLRMKDQCDKCREILSVDCSTNNPSQAKLRRE  
LDESLQVAERLTRKYNELLKSYQWKMLNTSSLLEQLNEQFNWVSRLANLTQGE  
DQYYLRVTTVASHTSDSDVPSGVTEVVVKLFSDPITVTVPVEVSRKNPKFMET  
VAEKALQEYRKKHREE

SEQ ID NO: 88

Human casein kinase 1, epsilon protein (GenBank# NM\_152221)

416 aa

MELRVGNKYRLGRKIGSGSFGDIYLGANIASGEEVAIKLECVKTKHPQLHIESKF  
YKMMQGGVGIPSIKWCGAEGDYNVMVMELLGPSLEDLFNFCSRKFSLKTVLLL  
ADQMISRIEYIHSKNFIHRDVKPDNFLMGLGKKGNLVYIIDFGLAKKYRDARTHQ

HIPYRENKNLTGTARYASINTHLGIEQSRDDLES LGYVLMYFNLGSLPWQGLKA  
ATKRQKYERISEKKMSTPIEVLCKGYPSEFSTYLNFCRSLRFDDKPDYSYLRQLFR  
NLFHRQGFSDYDYVFDWNMLKFGAARNPEDVDRERREHEREERMGQLRGSATR  
ALPPGPPTGATANRLRSAAEPVASTPASRIQPAGNTSPRAISRVDREKRVSMRLH  
RGAPANVSSDLTGRQEVSRIPASQTSVPFDHLGK

SEQ ID NO: 89

Human ferritin, heavy polypeptide 1 protein (GenBank# BC015946)

110 aa

MTTASTSQVRQNYHQDSEAAINRQINLELYASYVYLSMSYFDRDDVALKNFA  
KYFLHQSHEREHAEKLMKLQNQRGGRIFLQDIKKPDCDDWESGLNAMECALH  
LEKM

SEQ ID NO: 90

Human metargidin protein (GenBank# NM\_003815)

814 aa

MRLALLWALGLLGAGSPLPSWPLPNIGGTEEQQAESKAPREPLEPQVLQDDLPI  
SLKKVLQTSLEPLRIKLELDGDSHILELLQNRELVPGRPTLVWYQPDGTRVSE  
GHTLENCCYQGRVRGYAGSWVSICTCSGLRGLVLTPEPSTYLEQGPDLQGPPI  
ISRIQDLHLPGHTCALSWRESVHTQTPPEHPLGQRHIRRRRDVVTETKTVELVIVA  
DHSEAQKYRDFQHLLNRTLEVALLLDTFFRPLNVRVALVGLEAWTQRDLVEISP  
NPAVTLENFLHWRRRAHLLPRLPHDSAQLVTGTSFSGPTVGMAIQNSICSPDFSGG  
VNMDHSTSILGVASSIAHELGHSLGLDHDLPGNPCPCPGPAPAKTCIMEASTDFLP  
GLNFSNCSRRALEKALLDGMGSCLEFRLPSLPPMAAFCGNMFVEPGEQCDGFL  
DDCVDPCDSLTCQLRPGAQCASDGPCQCNCQLRPSGWQCRPTRGDCDLPEFCP  
GDSSQCPPDVSLGDGEPACAGGQAVCMHGRCASYAQQCQSLWGPGAQPAAPLCL  
QTANTRGNAFGSCGRNPSGSYVSCTPRDAICGQLQCQTGRTQPLLGSIRDLLWET  
IDVNGTELNCSSWVHLDLGSDVAQPLLTLPGTACGPGLVCIDHRCQRVDLLGAQE  
CRSKCHGHGVCDSNRHCYCEEGWAPPDCTTQLKATSSLTGLLSLLVLLVLM  
LGAGYWYRARLHQRLCQLKGPTCQYRAAQSGPSEPPGPPQRAALLARGTKSQGP  
AKPPPPRKPLPADPQGRCPSGDLPGPGAGIPPLVVPSPAPPPPTVSSLYL

SEQ ID NO: 91

Human unknown protein PHG-13 peptide 1 (GenBank# AK026351)

55 aa

MNLSFREFNQEKRVGGISWGPKGRLSGIFSTIQNQQSQKRGMSNSL  
KRTPQNS

SEQ ID NO: 92

Human unknown protein PHG-13 peptide 2 (GenBank# AK026351)

54 aa

MGNQRWHAKFNSGLRYPHCPHQASPALTVEPHGEEHVLERDPFVNCF  
VVFSSMN

SEQ ID NO: 93

Human unknown protein PHG-13 peptide 3 (GenBank# AK026351)

51 aa

MLCAQGAAGCQQHLSLNTISLCAEKTGNQRINITSPGWRTISCDFAAE  
FTH

SEQ ID NO: 94

Human unknown protein PHG-13 peptide 4 (GenBank# AK026351)

43 aa

MPPLIPHAAKRIGTLSGPGTVVMAISYFTHTRPFKVSLPQAIK

SEQ ID NO: 95

Human unknown protein PHG-13 peptide 5 (GenBank# AK026351)

39 aa

MVENIPESLPFGPQLMPPTLFSWLNSLKERFMCYCPVSQ

SEQ ID NO: 96

Human unknown protein PHG-13 peptide 6 (GenBank# AK026351)

36 aa

MSQCTSYPLIQKEEHFAQRKIKRSMNVIFYLLFSVG

SEQ ID NO: 97

Human unknown protein PHG-13 peptide 7 (GenBank# AK026351)

33 aa

MGSSLPIGFL LHTAGLSLYFKKKKKKKDKNCH

SEQ ID NO: 98

Human retinaldehyde binding protein 1 (GenBank# NM\_000326)

317 aa

MSEGVGTFRMVPEEEQELRAQLEQLTTKDHGPVFGPCSQLPRHTLQKAKDELNE  
REETREEAVRELQEMVQAQAASGEELAVAVAERVQEKDSGFFLRFIRARKFNVG  
RAYELLRGYVNFRLQYPELFDLSPEAVRCTIEAGYPGVLSSRDKYGRVVMLENI  
ENWQSQEITFDEILQAYCFILEKLLENEETQINGFCIENFKGFTMQQAASLRTSDL  
RKMVDMLQDSFPAKFKAHFHQPWYFTTTYNVVKPFLKSKLLERVVHGGDLS  
GFYQEIDENILPSDFGGTLPKYDGKAVAEQLFGPQAQAENTAF

SEQ ID NO: 99

Human actin, gamma 1, protein (GenBank# BC009848)

375 aa

MEEELAAALVIDNGSGMCKAGFAGDDAPRAVFPSIVGRPRHQGVMMVGMGQKDSY  
VGDEAQSKRGILTLKYPIEHGIVTNWDDMEKIWHHTFYNELRVAPEEHPVLLTE

APLNPKANREKMTQIMFETFNTPAMYVAIQAVLSLYASGRTTGIVMDSGDGVTH  
TVPIYEGYALPHAILRLDLAGRDLTDYLMKILTERGYSFTTTAEREIVRDIKEKLC  
YVALDFEQEMATAASSSSLEKSYELPDGQVITIGNERFRCPEALFQPSFLGMESCG  
IHETTFNSIMKCDVDIRKDLYANTVLSGGTTMYPGIADRMQKEITALAPSTMKIKI  
IAPPERKYSVWIGGSILASLSTFQQMWISKQEYDESGPSIVHRKCF

SEQ ID NO: 100

Human matrix metalloproteinase, membrane associated, protein (GenBank# X83535)

582 aa

MSPAPRPSRCLLLPLLTLGTALASLGSAQSSSFSPAWLQQYGYLPPGDLRHTQ  
RSPQSLSAIAAMQKFYGLQVTGKADATMKAMRRPRCGVPDKFGAEIKANVR  
RKRYAIQGLKWQHNEITFCIQNYTPKVGEYATYEAIRKAFRVWESATPLRFREVP  
YAYIREGHEKQADIMIFFAEGFHGDSTPFDGEGGFLAHAYFPGPNIGGDTHFDSA  
EPWTVRNEDLNGNDIFLVAVHELGHALGLEHSSDPSAIMAPFYQWMDTENFVLP  
DDDRRGIIQQLYGGESGFPTKMPPQPRRTTSRPSVPDKPKNPTYGPNICDGNFDTVA  
MLRGEMFVFKERWFWVRNNQVMDGYPMPIGQFWRGLPASINTAYERKDGKF  
VFFKGDKHWVFDEASLEPGYPKHIKELGRGLPTDKIDAALFWMPNGKTYFFRGN  
KYYRFNEELRAVDSEYPKNIKVWEGIPESPRGSFMGSDEVFTYFYKGNKYWKFN  
NQKLKVEPGYPKSALRDWMGCPSGGRPDGTEETEVIIEVDEEGGGAVSAAA  
VVLPLVLLLLLVLA VGLAVFFFRRHGTTPRRLLYCQRSLLDKV

SEQ ID NO: 101

Human SWI/SNF related/OSA-1 nuclear protein (GenBank# NM\_006015)

2285 aa

MAAQVAPAAASSLGNPPPPPPSELKKAEEQQQREEAGGEAAAAAAAAAERGEMKAA  
AGQESEGPVAVGPPQPLGKELQDGAESNGGGGGGGGAGSGGGPGAEPDLKNSNGN  
AGPRPALNNNLTEPPGGGGGGSSDGVGAPPHSAAAAALPPPAYGFGQPYGRSPSA  
VAAAAAAVFHQHGGQQSPGLAALQSGGGGGLEPYAGPQQNSHDHGFNPQHQY  
NSYYPNRSAYPPPAPAYALSSPRGGTPGSGAAAAAGSKPPSSSASASSSSSSFAQ  
QRFAMGGGGPSAAGGGTPQPTATPTLNQLLTSPSSARGYQGYPGGDYSGGPQ  
DGGAGKGPADMASQCWGAAAAAAAAAAAAASGGAQQRSHHAPMSPGSSGGGGQ  
PLARTPQPSSPMDQMGMKMRPQPYGGTNPYSQQQGGPPSGPQQGHGYPGQPYGSQ  
TPQRYPMTMQGRAQSAMGGLSYTQQIPYGGQQGPGSYGQQGQTPYNNQQSPHP  
QQQQPPYSQQPPSQTPHAQPSYQQQPQSQQPQLQSSQPPYSQQPSQPPHQQSPAP  
YPSQQSTTQQHPQSQPPYSQPQAQSPYQQQPPQPPAPSTLSQQAAYPQPQSQQSQ  
QTAYSQQRFPPPPQELSQDSFGSQASSAPSMTSSKGGQEDMNLSLQSRPSSLPDLS  
GSIDDLPMGTEGALSPGVSTSGISSQGEQSNPAQSPFSPHTSPHLPGIRGPSPPSPVG  
SPASVAQSRSGPLSPAAPVGNQMPPRPPSGQSDSIMHPSMNQSSIAQDRGYMQRN  
PQMPQYSSPQPGSALSPRQPSGGQIHTGMGSYQQNSMGSYGPPGGGQYGPQGGY  
PRQPNYNALPNANYPAGMAGGINPMGAGGQMHHGQPGIPPYGTLPGRMSHAS  
MGNRPYGPNNMANMPPQVSGMCPPPGGMNRKTQETAVAMHVAANSIQNRPPG  
YPNMNQGGMMGTGPPYGGGINSMAGMINPQGPYSMGGTMANNAGMAASPE  
MMGLGDVKLTPATKMNNKADGTPKTESKSKSSSSTTTNEKITKLYELGGEPER  
KMWVDRYLAFTEEKAMGMTNLPAVGRKPLDLYRLYVSVKEIGGLTQVNKNKK  
WRELATNLNVGTSSSAASSLKKQYIQCLYAFECKIERGEDPPPDIFAAADSKKSQ

PKIQPPSPAGSGSMQGPQTPQSTSSSMAEGGDLKPPTPASTPHSQIPPLPGMSRSNS  
VGIQDAFNDGSDSTFQKRNSMTPNPGYQPSMNTSDMMGRMSYEPNKDPYGSMR  
KAPGSDPFMSSGQGPNGGMGDPYSRAAGPGLGNVAMGPRQHYPYGGPYDRVR  
TEPGIGPEGNMSTGAPQPNLMPSNPDSGMYSRYPYPPQQQQQQQQRHDSYGNQF  
STQGTSPSGSPFPSQQTMYQQQQQNYKRPMDGTYGPPAKRHEGEMYSVPYSTG  
QGQPQQQQQLPPAQPPASQQQAAQPSPPQDVYNQYGNAYPATATAATERRPAG  
GPQNQFPFQFGRDRVSAPPGTNAQQNMPPQMMGGPIQASAEVAQQGTMWQGR  
NDMTYNYANRQSTGSAPQGPAYHGVNRTDEMLHTDQRANHEGSWPSHGTRQP  
PYGPSAPVPPMTRPPPSNYQPPPSMQNHIPQVSSPAPLPRPMENRTSPSKSPFLHSG  
MKMQKAGPPVPASHIAPAPVQPPMIRRDITFPPGSVEATQPVLKQRRRLTMKDIG  
TPEAWRVMMSLKSGLLAESTWALDTINILLYDDNSIMTFNLSQLPGLLELLVEYF  
RRCLIEIFGILKEYEVGDPGQRTLLDPGRFSKVSSPAPMEGGEEEEELLGPKLEEEE  
EEEVVENDEEIAFSKGDKPASENSEEKLISKFDKLPVKIVQKNDPFVVDKCDKLG  
RVQEFDSGLLHWRIJGGDTTEHIQTHFESKTELLPSRPHAPCPPAPRKHVTTAEG  
TPGTTDQEGPPPDGPPEKRITATMDDMLSTRSSTLTEDGAKSSEAIKESSKFPFGIS  
PAQSHRNKILEDEPHSKDETPLCTLLDWQDSLAKRCVCVSNTIRSLSFVPGNDFE  
MSKHPGLLLLILGKLILLHHKHPERKQAPLTYEKEEEEQDQGVSCNKVEWWDCLE  
EMLRENTLVTLANISGQLDLSPYPESICLPVLDGLLHWAVCPSEAEQDPFSTLGP  
NAVLSPQRLVLETLSKLSIQDNNVDLILATPPFSRLEKLYSTMVRFLSDRKNPVC  
EMAVVLLANLAQGDSLAARAIAVQKGSIGNLLGFLEDSLAATQFQQSQASLLHM  
QNPPFEPTSVDDMMRRAARALLALAKVDENHSEFTLYESRLDISVSPLMNSLVSQ  
VICDVLFLIGQS

SEQ ID NO: 102

Human unknown protein AMDP-3 peptide 1(GenBank# AK024103)

88 aa

MATQARQETCDNTKWN SHYARSCDHHQYHPQRSYKAKA  
HKGAPGGRWC VQG VGWHVCVGAHCHGASISKNSSREVC  
AEILACIPKAAH

SEQ ID NO: 103

Human unknown protein AMDP-3 peptide 2(GenBank# AK024103)

69 aa

MPYDSVRIERRMRCFKSKSQLLDSQVFKYGHPTPYLVLDY  
MGYEQGIETDKIVFTDTVYRFFFPFMQLFS

SEQ ID NO: 104

Human unknown protein AMDP-3 peptide 3(GenBank# AK024103)

65 aa

MCFNFKMLNSFQTWYLIYSPFLAFVEFQAECLTDCPRTL  
SFNLKQLRKGGQRRYKGKAAQNRSGE

SEQ ID NO: 105

Human unknown protein AMDP-3 peptide 4(GenBank# AK024103)

61 aa

MLGAVITTNITPRGVIKPRRTRGPLVAGGVCRGLGGTSVL  
VPTVTVQASARTQAGKSVLKY

SEQ ID NO: 106

Human unknown protein AMDP-3 peptide 5(GenBank# AK024103)

58 aa

MCNFFKYVFYSYGLLVSEPDLLTIFLYNNASHFLDSLVMC  
CMQELSSSSEGGLPLQAS

SEQ ID NO: 107

Human unknown protein AMDP-3 peptide 6(GenBank# AK024103)

55 aa

MLKKKNFFLVEMQSPVKRYEKASLSQRPGRQSTTRGSEV  
LMESCLSNEVLKRMPK

SEQ ID NO: 108

Human unknown protein AMDP-3 peptide 7(GenBank# AK024103)

50 aa

MLQIRKLLLGTCDTHSECDMVANGWPVLKAGSQHKGQR  
ALAAPLPTSEPG

SEQ ID NO: 109

Human unknown protein AMDP-3 peptide 8(GenBank# AK024103)

49 aa

MRHHLFYKLDYGFKWNTQGNIYKHQGKLSTASLFHLERG  
RFPNQTGFD P

SEQ ID NO: 110

Human unknown protein AMDP-3 peptide 9(GenBank# AK024103)

48 aa

MPVHSSLGNKSETPCQKKKKKMLLILSESKKETLTALNSG  
FIFLAVFG

SEQ ID NO: 111

Human unknown protein AMDP-3 peptide 10 (GenBank# AK024103)

48 aa

MRSWDL LFSPGLQNLIPVTKARKELYHKPSLSWHENWLP  
GSVYPINCE

SEQ ID NO: 112

Human unknown protein AMDP-3 peptide 11 (GenBank# AK024103)

45 aa

MIGHEASCHTPEIRVRLLLRTMCLVTYFSKIISLPGNQSSL  
VYLS

SEQ ID NO: 113

Human unknown protein AMDP-3 peptide 12 (GenBank# AK024103)

45 aa

MFIIIFIKVCVIFLSMYSIHMVCLSVSQTCLLYSFIIMLATS  
WIL

SEQ ID NO: 114

Human unknown protein AMDP-3 peptide 13 (GenBank# AK024103)

44 aa

MRTGCQAQCTPLTVNESELGFLYCFLCNMIAETHFKNSEA  
CHSC

SEQ ID NO: 115

Human unknown protein AMDP-3 peptide 14 (GenBank# AK024103)

40 aa

VMAYYSQGQVCPAQGVISGGFQTCTQFKDGGDRLCLYLVN  
PT

SEQ ID NO: 116

Human unknown protein AMDP-3 peptide 15 (GenBank# AK024103)

39 aa

MISAHCDLRLLGSSDSPASASRVAGITGMRHHARLILYF

SEQ ID NO: 117

Human unknown protein AMDP-3 peptide 16 (GenBank# AK024103)

39 aa

MEDFFLTALFFMAFSKRFKCSLFFKWGSLGRGKVCPHHL

SEQ ID NO:118

Human unknown protein AMDP-3 peptide 17 (GenBank# AK024103)

39 aa

M L E A L W N S P I P P P F Y I S L P T L A P M L L V P L Q C I P T Q G S I P

SEQ ID NO: 119

Human unknown protein AMDP-3 peptide 18 (GenBank# AK024103)

34 aa

M Y S T K M E P Y A W A L G I Q A S I S A Q T S L L E F L L M L A P

SEQ ID NO: 120

Human unknown protein AMDP-3 peptide 19 (GenBank# AK024103)

33 aa

M V S S P Q G G E A T H T M L K I N T K N K H K V R L V L H M C D

SEQ ID NO: 121

Human MT1-MMP exon 5 PCR product protein

53 aa

N D I F L V A V H E L G H A L G L E H S S D P S A I M A P F Y Q W M D T E N F V L P D D D R  
R G I Q Q L Y

SEQ ID NO: 122

Human MT1-MMP splice variant protein

260 aa

M S P A P R P S R C L L L P L L T L G T A L A S L G S A Q S S S F S P E A W L Q Q Y G Y L P P G D L R T H T Q  
R S P Q S L S A A I A A M Q K F Y G L Q V T G K A D A D T M K A M R R P R C G V P D K F G A E I K A N V R  
R K R Y A I Q G L K W Q H N E I T F C I Q N Y T P K V G E Y A T Y E A I R K A F R V W E S A T P L R F R E V P  
Y A Y I R E G H E K Q A D I M I F F A E G F H G D S T P F D G E G G F L A H A Y F P G P N I G G D T H F D S A  
E P W T V R N E D L N G N D I F L V A V H E L G H A L G L E H S S D P S A I M A P G

SEQ ID NO: 123

Human MT1-MMP exon 5 PCR product protein with D273N polymorphism

53 aa

N D I F L V A V H E L G H A L G L E H S S D P S A I M A P F Y Q W M D T E N F V L P N D D R  
R G I Q Q L Y

SEQ ID NO: 124

Human ABCR protein (GenBank# NM\_000350)

2273 aa

MGFVRQIQLLLWKNWTLRKRQKIRFVVELVWPLSLFLVLIWLRNANPLYSHHEC  
HFPNKAMPSAGMLPWLQGIFCNVNNPCFQSPTPGESPGIVSNYNNSILARVYRDF  
QELLMNAPESQHLGRIWTELHILSQFMDTLRTHPERIAGRIRIRDILKDEETLTLF  
LIKNIGLSDSVVYLLINSQVRPEQFAHGVPDLALKDIACSEALLERFIIFSQRRGAK  
TVRYALCSLSQGTQLQWIEDTLYANVDFKLFRLVPTLLDSRSQGINLRSWGILS  
DMSPRIQEFIHPSMQDLLWVTRPLMQNGGPETFTKLMGILSDLLCGYPEGGGSR  
VLSFNWYEDNNYKAFLGIDSTRKDPISYDRRTTSFCNALIQSLESNPLTKIAWR  
AAKPLLMGKILYTPDSPAARRILKNANSTFEELEHVRKLVKAWEEVGPQIWFYFF  
DNSTQNMNIRDITLGNPTVKDFLNRQLGEEGITAAILNFLYKGPRESQADDMAN  
FDWRDIFNITDRTLRLVNQYLECLVLDKFESYNDETQLTQRALSLEENMFWAG  
VVFPDMYPWTSSLPPHVKYKIRMDIDVVEKTNKIKDRYWDSGPRADPVEDFRYI  
WGGFAYLQDMVEQGITRSQVQAEAPVGIYLLQMPYPCFVDDSFMIILNRCFPIFM  
VLAWIYSVSMTVKISIVLEKELRLKETLKNQGVSNNAVIWCTWFLDSFSIMSMSIFL  
LTIFIMHGRILHYSDPFILFLFLLAFSTATIMLCFLLSTFFSKASLAAACSGVIYFTL  
YLPFILCFAWQDRMTAELKKAVSLLSPVAFGFGTEYLRVFEEQGLGLQWSNIGN  
SPTEGDEFSLLSMQMMLLDAACYGLLAWYLDQVFPGDYGTPLPWYFLLQESY  
WLSGEGCSTREERALEKTEPLTEETEDPEHPEGIHDSFFEREHPGWVPGVCVKNL  
VKIFEPGRPAVDRLNITFYENQITAFGLHNGAGKTTTSLTGLLPPTSGTVLVG  
GRDIETSLDAVRQSLGMCPQHNLFHHLTVAEHMLFYAQLKGKSQEEAQLMEEA  
MLEDTGHLHHRNEEAQDLSGGMQRKLSVAIAFVGDAKVVILDEPTSGVDPYSRR  
SIWDLKLYRSGRTIIMPTHMDEADHQGDRIIAAQGRLYCSGTPLFLKNCFGTG  
LYLTLVRKMKNIQSQRKSGEGTSCSSKGFSTTCPAHVDDLTPQVLDGDVNEL  
MDVVLHHVPEAKLVEICIGQELIFLLPNKNFKHRAVASLFRELEETLADLGLSSFGI  
SDTPLEEFLKVTEDSDSGPLFAGGAQQKRENVNPRHPCLGPREKAGQTPQDSNV  
CSPGAPAAHPEGQPPPECEPCGPQLNTGTQLVLQHVQALLVKRFQHTIRSHKDFL  
AQIVLPATFVFLALMLSIVILPFGEYPALTLHPWIYGQQYTFFSMDEPGSEQFTVL  
ADVLLNKPGFGNRCLKEGWLPEYPCGNSTPWKTPSVSPNITQLFQKQKWTQVNP  
SPSCRCSTREKLTMPECEGAGGLPPPQRTQRSTEILQDLTDRNISDFLVKTYPA  
LIRSSLKSKFWVNEQRYGGISIGGKLPVVPITGEALVGFLSDLGRIMNVSGGPITRE  
ASKEIPDFLKHLETEDNIKVWFNNKGWHALVSFLNVAHNAILRASLPKDRSPEEY  
GITVISQPLNLTKEQLSEITVLTTSDAVVAICVIFSMSFVPASFVLYLIQERVNKS  
KHLQFISGVSPPTYWVTNFLWDIMNYSVSAGLVVGIFIGFQKKAYTSPENLPALV  
ALLLYGWAVIPMMYPASFLFDVPSTAYVALSCANLFIGINSSAITFILELFDNNR  
TLLRFNAVLRKLLIVFPHFCLGRGLIDLALSQAVTDVYARFGEEHSANPFHWDLI  
GKNLFAMVVEGVVYFLLTLLVQRHFFLSQWIAEPTKEPIVDEDDDDVAEERQRIIT  
GGNKTDILRLHELTKIYLGTSPPAVDRLCVGVRPGECFGLLGVNAGKTTTFKM  
LTGDTTVTSGDATVAGKSILTNISEVHQNMGYCPQFIDAIDELLTGREHLYLYARL  
RGVPAEEIEKVANWSIKSLGLTVYADCLAGTYSGGNKRKLSTAIALIGCPPLVLL  
DEPTTGMDPQARRMLWNVIVSIIRKGRAVVLTSHEMEECEALCTRLAIMVKGAF  
RCMGTIQHLKSKFGDGYIVTMKIKSPKDDLLPDLNPVEQFFQGNFPGSVQRRERHY  
NMLQFQVSSSSSLARIFQLLLSHKDSLLIEEYSVTQTTLQVFNFAKQQTESHDLF  
LHPRAAGASRQAQD

SEQ ID NO: 125

Human apolipoprotein E protein (GenBank# NM\_000041)

317 aa

MKVLWAALLVTFLAGCQAKVEQAVETEPEPELRQQTEWQSGQRWELALGRFW  
DYLRWVQTLSEQVQEELLSSQVTQELRALMDETMKELKAYKSELEEQLTPVAEE  
TRARLSKELQAAQARLGADMEDVCGRLVQYRGEVQAMLGQSTEELRVRASHL  
RKLRKRLLRDADDLQKRLAVYQAGAREGAERGLSAIRERLGPLVEQGRVRAAT  
VGSLAQPLQERAQAWGERLRARMEEMGSRTDRDLDEVKEQVAEVRAKLEEQ  
AQQIRLQAEAFQARLKSWFEPLVEDMQRQWAGLVEKVQAAVGTSAAPVPSDN  
H

SEQ ID NO: 126

Human C-C chemokine receptor-2 (Ccr-2) protein (GenBank# NM\_000647)

374 aa

MLSTSRSRFIRNTNESGEEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSLVFIFG  
FVGNMLVVLILINCKKLKCLTDIYLLNLAISDLLFLITLPLWAHSAANEVVFNGA  
MCKLFTGLYHIGYFGGIFFIILLTIDRYLAIVHAVFALKARTVTFGVVTSVITWLV  
AVFASVPGIIFTKCQKEDSVYVCGPYFPRGWNNFHTIMRNILGLVLPLLIMVICYS  
GILKTLLRCRNEKKRHRAVRVIFTIMIVYFLFWTPYNIVILLNTFQEFFGLSNCEST  
SQLDQATQVTETLGMTHCCINPIIYAFVGEKFRSLFHIALGCRIAPLQKPVCGGPG  
VRPGKNVKVTTQGLLDGRGKGKSGRAPEASLQDKEGA

SEQ ID NO: 127

Human cystatin C protein (GenBank# NM\_000099)

146 aa

MAGPLRAPLLLLAILAVALAVSPAAGSSPGKPPRLVGGPMDASVEEEGVRRALD  
FAVGEYNKASNDMYHSRALQVVRARKQIVAGVNYFLDVELGRTTCTKTQPNLD  
NCPFHDQPHLKRKAFCSFQIYAVPWQGTMTLSKSTCQDA

SEQ ID NO: 128

Human hemicentin/FIBL-6 protein (GenBank# NM\_031935)

5622 aa

MISWEVVHTVFLFALLYSSLAQDASPQSEIRAEIPEGASTLAFVFDVTGSMYDD  
LVQVIEGASKILETSLKRPKRPLFNFALVPFHDPEIGPVTITTDPKKFQYELRELYV  
QGGGDCPEMSIGAIIKIALEISLPGSFIYVFTDARSKDYRLTHEVLQLIQKQSQVV  
FVLTGDCDDRTHIGYKVYEEIASTSSGQVFHLDKKQVNEVLKWVEEAVQASKV  
HLLSTDHLEQAVNTWRIPFDPSLKEVTVSLSGSPMIEIRNPLGKLIKKGFLHEL  
LNIHNSAKVVNVKEPEAGMWTVKTSSSGRHSVRITGLSTIDFRAGFSRKPTLDFK  
KTVSRPVQGIPTYVLLNTSGISTPARIDLLELLSISGSSLKTIPVKYYPHRKPYPYGIW  
NISDFVPPNEAFFLKVTGYDKDDYLFQRVSSVSFSSIVPDAPKVTMPEKTPGYL  
QPGQIPCSVDSLPLFTLSFVRNGVTLGVDQYLKESASVNLDIAKVTLSDGEFYECI  
AVSSAGTGRAQTFFDVSEPPPIQVPNNVTVTPGERAVLTCLISAVDYNLTWQR

NDRDVRLAEPARIRTLANLSLELKS VKFNDAGEYHCMVSSEGGSSAASVFLTVQ  
EPPKVTVM PKNQSF TGGSEVSIMCSATGYPKPKIAWTVNDMFIVGSHRYRMTSD  
GTLFIKNAAPKDAGIYGCLASNSAGTDKQNSTLRYIEAPKLMVVQSELLVALGDI  
TVMECKTSGIPPPQVKWFKGDLELRPSTFLIIDPLLGLLKIQETQDLDAGDYTCVA  
INEAGRATGKITLDVGSPPVFIQEPADVSM EIGSNVTLPCYVQGYPEPTIKWRRLD  
NMPIFSRPFSVSSISQLRTGALFILNLWASDKGT YICEAENQFGKIQSETTVTVTGL  
VAPLIGISPSVANVIEGQQLTLPCTLLAGNP IERRWIKNSAMLLQNPYITVRS DGS  
LHIERVQLQDGG EYTCVASNVAGTNNKTT SVVVHVLPTIQHGQQILSTIEGIPVTL  
PCKASGNPKPSVIW SKKGELISTSSAKFSAGADGSLYVVSPGGEESGEYVCTATN  
TAGYAKRKVQLTVYVRPRVFGDQRGLS QDKPVEISVLAGEEVTLPCEVKS LPPPI  
ITWAKETQLISPFSPRHTFLPSGSMKITETRTSDSGMYLCVATNIAGNVTQAVKLN  
VHVPPKIQRGPKHLKVQVGQRVDIPCNAQGTPLPVITWSKGGSTMLVDGEHHVS  
NPDGTLSDIDQATPSDAGIYTCVATNIAGTDETEITLHVQEPPPTVEDLEPPYNTTFQE  
RVANQRIEFP CPAKGT PKPTIKWLHNGREL TGREPGISILEDGTL LVIASVTPYDN  
GEYICVAVNEAGTTERKYNLKVHVPPVIKDKEQVTNVS VLLNQLTNLFCEVEGT  
PSPIIMWYKDNVQVTESSTIQTVNNGKILKLFRATPEDAGRYSC KAINIAGTSQKY  
FNIDVLVPPTIIGTNFPNEVSVVLNRDVALECQVKGT PFPDIHWFKDGKPLFLGDP  
NVELLDRGQVLHLKNARRNDKG RYQCTVSNAAGKQAKDIKLT IYIPPSIKGGNV  
TTDISVLINSLIKLECETRGLPMPAITWYKDGQPI MSSSQALYIDKGQYLHIPRAQV  
SDSATYTCHVANVAGTAEKSFHVDVYVPPMIEGNLATPLNKQV VIAHSLTLECK  
AAGNPSPILT WLKDGVPVKANDNIRIEAGGKKLEIMSAQEIDRGQYICVATSVAG  
EKEIKYEVDVLVPPAIEGGDETSYFIVMVNNLLELDCHVTGSPPTIMWLKDGQLI  
DERDGFKILLNGRKL VIAQAQVSNTGLYRCMAANTAGD HKKEFEVTVHVPPPTIK  
SSGLSERVVVKYKPVALQCIANGIPNPSITWLKDDQPVNTAQGNLKIQSSGRVLQ  
IAKTLL EDAGRYTCVATNAAGETQQHIQLHVHEPPSLEDAGKMLNETVLVSNPV  
QLECKAAGNPVPVITWYKDNRLLSGSTSMTFLNRGQIIDIESAQISDAGIYKCVAI  
NSAGATELFYSLQVHVAPSISGSNNMVA VVVNNPVRLECEARGIPAPSLTWLKD  
GSPVSSFSNGLQVLSGGRILAL TSAQISDTGRYTCVAVNAAGEKQRDIDL RVYVP  
PNIMGEEQNVSVLISQAVELL CQSDAIPPTLTWLKDGHP LLKKPGLSISENRSVL  
KIEDAQVQDTGRYTCEATNVAGKTEKKNYNVNIWVPPNIGGSDEL TQLTVIEGN  
LISLLCESSGIPPPNLIWKKKGSPVLTDSMGRVRIIAEKSDAALYSCVASN VAGTA  
KKEYNLQVYIRPTITNSGSHPTEIIVTRGKSISLECEVQGIPPTVTWMKDGHPLIK  
AKGVEILDEGHILQLKNIHVSDTGRYVCVAVNVAGMTDKKYDLSVHAPPSIIGN  
HRSPENISVVEKNSVSLTCEASGIPLPSITWFKDGWPVSLSNSVRILSGGRMLRLM  
QTTMEDAGQYTCVVRNAAGEERKIFGLSVLVPPHIVGENTLEDVKVKEKQSVTL  
TCEVTGNPVPEITWHKDGQPLQEDEAHHIISGGRFLQITNVQVPHTGRYTCLASSP  
AGHKSRFSFSLNVFVSPTIAGVGSDGNPEDVTVILNSPTSLVCEAYSYP PATITWFK  
DGTPLESNRNIRILPGGRTLQILNAQEDNAGRYSCVATNEAGEMIKHYEVKVYIP  
PIINKGDLWGPGLSPKEVKIKVNNTLTLECEAYAIP SASLSWYKDGOPLKSDDHV  
NIAANGHTLQIKEAQISDTGRYTCVASNIAGEDELD FDNIVQVPPSFQKLWEIGN  
MLDTGRNGEAKDVIINNPISLYCETNAAPPPTLTWYKDGHP LTSSDKVLILPGGR  
VLQIPRAKVEDAGRYTCVAVNEAGEDSLQYDVRVLVPPIIKGANS DLPEEVTVLV  
NKSALIECLSSGSPAPRNSWQKDGOPLLEDDH HKFLSNGRILQILNTQITDIGRYV  
CVAENTAGSAKKYFNLNVHVPPSVIGPKSENLT VVVNNFISLTCEVSGFPPPDLS  
WLKNKLNTNTLIVPGGRTLQIRAKVSDGGEYTCIAINQAGESKKKFS LTVYVPPS  
IKDHDSESLSVNVNREGTSVSLECESNAVPPPVITWYKNGRMITESTHVEILADG

QMLHIKKAEVSDTGQYVCRAINVAGRDDKNFHLNVYVPPSIEGPEREVIVETISN  
PVTLTCDATGIPPPTIAWLKNHKRIENSDDSLEVRILSGGSKLQIARSQHSDSGNYT  
CIASNMEGKAQKYYFLSIQVPPSVAGAEIPSDVSVLLGENVELVCNANGIPTPLIQ  
WLKDGKPIASGETERIRVSANGSTLNIYGALTSDTGKYTCVATNPAGEEDRIFNL  
NVYVTPTIRGNKDEAEKMLTLVDTSINIECRATGTPPPQINWLKNGLPLPLSSHIR  
LLAAGQVIRIVRAQVSDVAVYTCVASNRAGVDNKHYNLQVFAPPNMDNSMGTE  
EITVLKGSSTSMACITDGTAPASMAWLRDGOPLGLDAHLTVSTHGMVLQLLKAE  
TEDSGKYTCIASNEAGEVSKHFILKVLEPPHINGSEEHEEISVIVNNPLELTCIASGI  
PAPKMTWMKDGRLPQTDQVQTLGGGEVLRISTAQVEDTGRYTCLASPAGDD  
DKEYLVRVHVPPNIAGTDEPRDITVLRNRQVTLECKSDAVPPPVTWLRNGERLQ  
ATPRVRILSGGRYLQINNADLGDTANYTCVASNIAGKTTREFILTVNVPPNIKGGP  
QSLVILLNKSTVLECIAEGVPTPRITWRKDGAVLAGNHARYSILENGFLHIQSAHV  
TDTGRYLCMATNAAGTDRRRIDLQVHVPPSIAPGPTNMTVIVNVQTTLACEATGI  
PKPSINWRKNGHLLNVDQNNQNSYRLSSGSLVIISPSVDDTATYECTVTNGAGDD  
KRTVDLTVQVPPSIADPTDFLVTKHAPAVITCTASGVPPFSIHWTNGIRLLPRG  
DGYRILSSGAIEILATQLNHAGRYTCVARNAAGSAHRHVTLVHVEPPVIQPPSE  
LHVILNNPILLPCEATGTPSPFITWQKEGINVNTSGRNHAVLPSGGLQISRAVRED  
AGTYMCVAQNPAGTALGKIKLVQVPPVISPHLKEYVIAVDKPITLSCEADGLPP  
PDITWHKDGRAIVESIRQVRLSSGSLQIAFVQPGDAGHYTCMAANVAGSSSTSTK  
LTVHVPPRIRSTEGHYTVNENSQAILPCVADGIPTAINWKKDNVLLANLLGKYT  
AEPYGELILENVVLEDSGFYTCVANNAAGEDTHTVSLTVHVLPTFTELPGDVSLN  
KGEQLRLSCKATGIPLPKLTWTFNNNIIPAHFDSVNGHSELVIERVSKEDSGTYVC  
TAENSVGFVKAIGFVYVKEPPVFKGDYPSNWIEPLGGNAILNCEVKGDPTPTIQW  
NRKGVDIEISHRIRQLGNGSLAIYGTVNEDAGDYTCVATNEAGVVERSMSLTLOS  
PPIITLEPVETVINAGGKIILNCQATGEPQPTITWSRQGHSSWDDRNVNLSNNSLY  
IADAQKEDTSEFECVARNLMGSVLVRVPVIVQVHGGFSQWSAWRACSVTCGKG  
IQKRSRLCNQPLPANGGKPCQGSLEMRNCQNKPCPVDGSWSEWSLWEECTRS  
CGRGNQTRTRTCNNPSVQHGGRPCEGNAVEIIMCNIRPCPVHGAWSAWQPWGT  
CSESCGKGTQTRARLCNNPPPAFGGSYCDGAETQMQVCNERNCPHKGWATWA  
SWSACSVSCGGGARQRTRGCSDPVPQYGGRKCEGSDVQSDFCNSDPCPTHGNW  
SPWSGWGTCSTRTCNGGQMRRYRTCDNPPPSNGGRACGGPDSQIQRcntDMCPV  
DGSWGSWHSWSQCSASCGGGEKTRKRLCDHPVPVKGGRPCPGDttQVTRCNV  
QACPGGPQRARGSVIGNINDVEFGIAFLNATITDSPNSDTRIIRAKITNVPRSLGSA  
MRKIVSILNPIYWTTAKEIGEAVNGFTLTNAVFKRETQVEFATGEILQMSHIARGL  
DSDGSLLLDIVVSGYVLQLQSPA EVT VKDYTEDYIQTGPGQLYAYSTRLFTIDGIS  
IPYTNHTVFDYDQAQGRMPFLVETLHASSVESDYNQIEETLGFKIHASISKGDRS  
NQCPSGFTLDSVGPFCADEDECAAGNPCSHSCHNAMGTYYCSCP KGLTIAADGR  
TCQDIDECALGRHTCHAGQDCDNTIGSYRCVVRCGSGFRRTSDGLSCQDINECQ  
ESSPCHQRCFNAIGSFHCGCEPGYQLKGRKCMDVNECRQNVCRPDQHCKNTRG  
GYKCIDLCPNGMTKAENGTCIDIDECKDGTHQCRYNQICENTRGSYRCVCPRGY  
RSQGVGRPCMDINECEQVPKPCA HQCSNTPGSFKCICPPGQHLLGDGKSCAGLER  
LPNYGTQYSSYNLARFSPVRNNYQPQQHYRQYSHLYSSYSEYRNSRTSLSRTRRT  
IRKTCPEGSEASHDTCVDIDECENTDACQHECKNTFGSYQCICPPGYQLTHNGKT  
CQDIDECLEQNVHCGPNRMCFNMGRGSYQCIDTPCPPNYQRDPVSGFCLKNCPN  
DLECALSPYALEYKLVS L PFGIATNQDLIRLVAYTQDGMHPRTTFLMVDEEQT

VPFALRDENLKGVVYTTRPLREAETYRMRVRASSYSANGTIEYQTTFFIVYIAVSA  
YPY

SEQ ID NO: 129

Human manganese superoxide dismutase 2 protein (GenBank# NM\_000636)

222 aa

MLSRAVCGTSRQLPPVLGYLGSRQKHSLPDLPYDYGALPHINAQIMQLHHSKH  
HAAYVNNLNVTEEKYQEALAKGDVTAQIALQPALKFNGGGHINHSIFWTNLSPN  
GGGEPKGELLEAIKLDFGSFDKFKEKLTAA SVGVQSGWGWLGFNKERGHLQIA  
ACPNQDPLQGTTGLIPLLIDVWEHAYYLQYKNVRPDYKAIWNVINWENVTER  
YMACKK

SEQ ID NO: 130

Human C-C chemokine ligand 2 (Ccl-2)/monocyte chemoattractant protein 1 (GenBank#  
NM\_002982)

99 aa

MKVSAALLCLLLIAATFIPQGLAQPDAINAPVTCCYNFTNRKISVQRLASYRRITS  
SKCPKEAVIFKTIVAKEICADPKQKWVQDSMDHLDKQTQTPKT

SEQ ID NO: 131

Human paraoxonase 1 protein (GenBank# NM\_000446)

355 aa

MAKLIALTLLGMGLALFRNHQSSYQTRLNALREVQPVELPNCNLVKGIETGSED  
LEILPNGLAFISSGLKYPGIKSFNPNSPGKILLMDLNEEDPTVLELGITGSKFDVSSF  
NPHGISTFTDEDNAMYLLVVNHPDAKSTVELFKFQEEEEKSLLHLKTIRHKLLPNL  
NDIVAVGPEHFYGTNDHYFLDPYLQSWEMYLG LAWSYVVYYPSEVRVVAEGF  
DFANGINISPDGKYVYIAELLAHKIHVYEK HANWTLTPLKSLDFNTLVDNISVDP  
ETGDLWVGCHPNGMKIFFYDSENPPASEVLRIQNILTEEPKVTQVYAENGTVLQG  
STVASVYKGKLLIGTVFHKALYCEL